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(54) Title: GENE THERAPY FOR INHIBITION OF ANGIOGENESIS

(57) Abstract

The present invention relates to methods of gene therapy for inhibiting angiogenesis associated with solid tumor growth, tumor metastasis, inflammation, psoriasis, rheumatoid arthritis, hemangiomas, diabetic retinopathy, angiofibromas, and macular degeneration. Gene therapy methodology is disclosed for inhibition of primary tumor growth and metastasis by gene transfer of a nucleotide sequence encoding a soluble form of a VEGF tyrosine kinase receptor to a mammalian host. The transferred nucleotide sequence transcribes mRNA and a soluble receptor protein which binds to VEGF in extracellular regions adjacent to the primary tumor and vascular endothelial cells. Formation of a sVEGF-R/VEGF complex will prevent binding of VEGF to the KDR and FLT-1 tyrosine kinase receptors, antagonizing transduction of the normal intracellular signals associated with vascular endothelial cell-induced tumor angiogenesis. In addition, expression of a soluble receptor tyrosine kinase may also impart a therapeutic effect by binding either with or without VEGFs to form non-functional heterodimers with full-length VEGF-specific tyrosine kinase receptors and thereby inhibiting the mitogenic and angiogenic activities of VEGFs.

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TITLE OF THE INVENTION
GENE THERAPY FOR INHIBITION OF ANGIOGENESIS

CROSS-REFERENCE TO RELATED APPLICATIONS

This application is a continuation-in-part application of U.S. Provisional Application Serial No. 60/026,641, filed September 24, 1996.

STATEMENT REGARDING FEDERALLY SPONSORED RESEARCH

Not applicable.

REFERENCE TO MICROFICHE APPENDIX

Not applicable.

FIELD OF THE INVENTION

The present invention relates to methods of gene therapy for inhibiting angiogenesis associated with tumor growth, inflammation, psoriasis, rheumatoid arthritis, hemangiomas, diabetic retinopathy, angiofibromas, and macular degeneration.

This invention also relates to animal models useful in the investigation of gene therapy-mediated inhibition of angiogenesis. The invention also relates to recombinant vectors which are useful in the disclosed gene therapy methods.

BACKGROUND OF THE INVENTION

Vascular endothelial cells form a luminal non-thrombogenic monolayer throughout the vascular system. Mitogens promote embryonic vascular development, growth, repair and angiogenesis in these cells. Angiogenesis involves the proteolytic degradation of the basement membrane on which endothelial cells reside followed by the subsequent chemotactic migration and mitosis of these cells to support sustained growth of a new capillary shoot. One class of mitogens selective for vascular endothelial cells include vascular endothelial growth factor (referred to as VEGF or VEGF-A) and the homologues placenta growth factor (PlGF), VEGF-B and VEGF-C.

Human VEGF exists as a glycosylated homodimer in one of five mature processed forms containing 206, 189, 165, 145 and 121 amino acids, the most prevalent being the 165 amino acid form.

U.S. Patent No. 5,240,848 discloses the nucleotide and amino acid sequence encoding the 189 amino acid form of human VEGF.

U.S. Patent No. 5,332,671 discloses the nucleotide and amino acid sequence encoding the 165 amino acid form of human VEGF.

Charnock-Jones et al (1993, *Biol. Reproduction* 48: 1120-1128) discloses the VEGF145 splice variant m RNA.

U.S. Patent No. 5,194,596 discloses the nucleotide and amino acid sequence encoding the 121 amino acid form of human VEGF.

The 206 amino acid and 189 amino acid forms of human VEGF each contain a highly basic 24-amino acid insert that promotes tight binding to heparin, and presumably, heparin proteoglycans on cellular surfaces and within extracellular matrices (Ferrara, et al., 1991, *J. Cell. Biochem.* 47: 211-218). The VEGF165 form binds heparin to a lesser extent while VEGF121 does not bind heparin.

Human PlGF is also a glycosylated homodimer which shares 46% homology with VEGF at the protein level. Differential

splicing of human PlGF mRNA leads to either a 170 amino acid or 149 amino acid precursor, which are proteolytically processed to mature forms of 152 or 131 amino acids in length, respectively (Maglione, et al., 1993, *Oncogene* 8: 925-931; Hauser and Weich, 1993, *Growth Factors* 9: 259-268).

VEGF-B was recently isolated and characterized (Olofsson, et al., 1996, *Proc. Natl. Acad. Sci.* 93: 2576-2581; Grimmond et al., 1996, *Genome Research* 6: 124-131). The full length human cDNAs encode 188 and 207 amino acid precursors wherein the NH_2 terminal portions are proteolytically processed to mature forms 167 and 186 amino acids in length. Human VEGF-B expression was found predominantly in heart and skeletal muscle as a disulfide-linked homodimer. However, human VEGF-B may also form a heterodimer with VEGF (*id.* @ 2580).

VEGF-C has also recently been isolated and characterized (Joukov, et al., 1996, *EMBO J.* 15: 290-298). A cDNA encoding VEGF-C was obtained from a human prostatic adenocarcinoma cell line. A 32 kDa precursor protein is proteolytically processed to generate the mature 23 kDa form, which binds the receptor tyrosine kinase, Flt-4.

VEGF-D was identified in an EST library, the full-length coding region was cloned and recognized to be most homologous to VEGF-C among the VEGF family amino acid sequences (Yamada, et al., 1997, Genomics 42:483-488). The human VEGF-D mRNA was shown to be expressed in lung and muscle.

VEGF and its homologues impart activity by binding to vascular endothelial cell plasma membrane-spanning tyrosine kinase receptors which then activates signal transduction and cellular signals. The Flt receptor family is a major tyrosine kinase receptor which binds VEGF with high affinity. At present the flt receptor family includes flt-1 (Shibuya, et al., 1990, *Oncogene* 5: 519-524), KDR/flk-1(Terman, et al., 1991, *Oncogene* 6: 1677-1683; Terman, et al., 1992, *Biochem. Biophys. Res. Commun.* 187: 1579-1586), and flt-4 (Pajusola, et al., 1992, *Cancer Res.* 52: 5738-5743).

The involvement of VEGF in promoting tumor angiogenesis has spawned studies investigating possible antagonists of the process.

Both polyclonal (Kondo, et al., 1993, *Biochem. Biophys. Res..Commun.* 194: 1234-1241) and monoclonal (Kim, et al., 1992, *Growth Factors* 7: 53-64; Kim, et al., 1993, *Nature* 362: 841-844) antibodies raised against VEGF have been shown to suppress VEGF activity *in vivo*. Anti-VEGF antibody strategies to interdict angiogenesis and its attendant tumor are also addressed in Kim et al. (1993, *Nature* 362: 841-844) and Asano et al. (1995, *Cancer Research* 55: 5296-5301).

Kendall and Thomas (1993, *Proc. Natl. Acad. Sci.* 90: 10705-10709) isolated and characterized a cDNA encoding a secreted soluble form of flt-1 from cultured human umbilical vein endothelial cells (HUVEC). The recombinant version of this protein was purified by binding to immobilized heparin. Isolated soluble flt-1 was shown to inhibit VEGF activity *in vitro*. No suggestion regarding gene transfer protocols were disclosed.

Millauer et al. (1994, *Nature* 367: 576-579) disclose *in vivo* inhibition of tumor angiogenesis by expression of an artificially generated flk-1 mutant in which the intracellular kinase domain but not the membrane-spanning anchor was deleted. The authors do not forward any teaching or suggestion that a soluble form of a VEGF tyrosine kinase receptor would be useful in gene therapy applications.

Neovascularization of malignant tumors is an integral process contributing to solid tumor growth and neoplastic progression (Kondo et al., 1993, *Biochemical & Biophysical Research Communications* 194: 1234-1241; Carrau et al., 1995, *Invasion & Metastasis* 15: 197-202). In this context, several studies have demonstrated a positive correlation between neovascularization in malignant tumors and poor clinical outcomes (Volm et al., 1996, *Anticancer Research* 16: 213-217; Toi et al., 1994, Japanese Journal of Cancer Research 85: 1045-1049; Shpitzer et al., 1996, *Archives of Otolaryngology -- Head & Neck Surgery*; 122: 865-868; Staibano et al., 1996, *Human Pathology* 27: 695-700; Giatromanolaki et al., 1996, *J. of Pathology* 179: 80-88). While the angiogenic process has several mediators, it appears that vascular end thelial growth factor (VEGF) may be a critical growth factor with respect to initiating the cascade of

events stimulating new blood vessel formation in several tumor types (Toi et al., 1996, *Cancer* 77: 1101-1106; Maeda et al., 1996, *Cancer* 77: 858-63; Anan et al., 1996, *Surgery* 119: 333-339).

Aiello et al. (1995, Proc. Natl. Acad. Sci. USA 92:10457-10461) disclose genetically engineered chimeric extracellular VEGF receptors to block angiogenesis in non-malignant cells.

Despite recent advances in identifying genes encoding ligands and receptors involved in angiogenesis, no gene therapy application has been forwarded which overcomes the deleterious effect this process has in promoting primary tumor growth and subsequent metastasis. The present invention addresses and meets this need.

SUMMARY OF THE INVENTION

The present invention relates to methods of gene therapy for inhibiting VEGF-induced angiogenesis associated with diseases and disorders including, but not limited to, solid tumor growth, tumor metastasis, inflammation, psoriasis, rheumatoid arthritis, hemangiomas, angiofibromas, diabetic retinopathy, and macular degeneration. These disorders are related in that VEGF acts as a mitogen to stimulate local angiogenesis from vascular endothelial cells which in turns exacerbates the condition.

The present invention relates to gene transfer of a DNA vector and concomitant *in vivo* expression of a soluble form of a tyrosine receptor kinase (sVEGF-R) within the mammalian host which binds VEGF or a VEGF homologue in and around the localized site of the disorder. The formation of a sVEGF-R/VEGF complex will inhibit binding of VEGF to the FLT-1 and KDR tyrosine kinase receptors spanning the vascular endothelial cell membrane, thus preventing initiation of the signal transduction stimulating angiogenesis. In addition, expression of sVEGF-R may also impart a therapeutic effect by binding to membrane associated VEGF-Rs. VEGF-Rs are thought to be dimerized by binding dimeric VEGF ligand which in turn allows the receptor intracellular tyrosine kinase domains to transphosphorylate each other generating phosphorylated tyrosine residues that facilitate

the subsequent binding and activation of downstream signal transduction proteins. sVEGF-Rs can form heterodimers with full-length VEGF-Rs that, because the sVEGF-Rs are devoid of an intracellular tyrosine kinase region, prevent receptor tyrosine kinase domain transphosphorylation, the initiation of signal transduction and thus VEGF-induced mitogenesis and angiogenesis in a dominant negative manner.

A nucleotide sequence encoding a sVEGF-R for inclusion in a gene therapy vector of the present invention may be chosen from a group of genes encoding tyrosine kinase receptors, namely from the group consisting of s*flt*-1, *flt*-1, KDR (also denoted flk-1), and *flt*-4. The resulting DNA fragment encodes a protein or protein fragment which binds VEGF and/or KDR/flk-1 and inhibits formation of a wild-type, functional VEGF-R/VEGF complex.

A preferred application of the present invention relates to promoting inhibition of solid tumor angiogenesis and metastasis by utilizing the disclosed gene therapy methodology. In particular, methods are disclosed for inhibition of primary tumor growth and metastasis by gene transfer of a nucleotide sequence encoding sVEGF-R to a mammalian host. The transferred nucleotide sequence transcribes mRNA and expresses sVEGF-R such that sVEGF-R binds to VEGF in extracellular regions adjacent to the primary tumor and vascular endothelial cells. Formation of a sVEGF-R/VEGF complex will prevent binding of VEGF to the KDR and FLT-1 tyrosine kinase receptors, antagonizing transduction of the normal intracellular signals associated with vascular endothelial cell-induced tumor angiogenesis. In addition, expression of sFLT-1 may also impart a therapeutic effect by binding either with or without VEGFs to form non-functional heterodimers with full-length VEGF-Rs and thereby inhibiting the mitogenic and angiogenic activities of VEGFs.

In a particular embodiment of the present invention a truncated version of a soluble or transmembrane form of FLT-1 (Shibuya, et al., 1990, *Oncogene* 5: 519-524) is utilized in gen therapy protocols. It will be within th purview of the skilled artisan to generate

a sVEGF-R or VEGF-RTMI construct expressing a truncated FLT-1 protein which binds to VEGF, a VEGF homologue and/or dimerizes with a full-length VEGF-R inhibiting its activation on the surface plasma membrane of vascular endothelial cells (Figure 1). Such a construct may be generated by recombinant DNA techniques known in the art using a DNA fragment encoding a partial or complete amino acid sequence of a FLT receptor. Using recombinant DNA techniques, DNA molecules are constructed which encode at least a portion of the VEGF receptor capable of binding VEGF without stimulating either mitogenesis or angiogenesis. Standard recombinant DNA techniques are used such as those found in Maniatis, et al. (1982, Molecular Cloning: A Laboratory Manual; Cold Spring Harbor Laboratory, Cold Spring Harbor, New York).

In another embodiment of the present invention a mutated version of FLT-1 is generated which is defective in protein kinase activity, namely a FLT-1 protein mutated at or around one or more known active sites for protein kinase activity. A *flt*-1 construction will express the extracellular domain, transmembrane domain and the mutated portion of the intracellular domain such that the resulting protein at least substantially inhibits related intracellular protein kinase activity.

In a preferred embodiment of the present invention, a naturally expressed alternatively spliced DNA encoding a soluble form of FLT-1 (Kendall and Thomas, 1993, *Proc. Natl. Acad. Sci.* 90: 10705-10709; U.S. Application Serial No. 08/232,538, hereby incorporated by reference; described herein as sVEGF-RI or sFLT-1 and listed as SEQ ID NO:1 (nucleotide sequence) and SEQ NO ID:2 (amino acid sequence) is the template for constructing a gene therapy vector wherein either expressed sFLT-1 or a biologically active truncated form binds VEGF and inhibits complex formation, dimerization and activation of full-length VEGF-Rs, and hence, pathological angiogenesis.

The present invention relates to both viral and non-viral recombinant vectors for delivery to the target hosts. To this end, a preferred non-viral recombinant plasmid described herein is

pcDNA3/sflt-1. An especially preferred recombinant plasmid of the present invention is pcDNAIAsFLT-1, as decribed in Example Section 5.

A recombinant adenovirus (Ad) system is preferred for delivery and prolonged expression within target cells proximal to a solid tumor. A particularly useful adenovirus system used in the present invention is described in Example 4.

Any sVEGF-R construct, including but in no way limited to sVEGF-RI and biologically active truncated forms, may be delivered to the mammalian host using a vector or other delivery vehicle. DNA delivery vehicles can include viral vectors such as adenoviruses, adeno-associated viruses, and retroviral vectors. See, for example: Chu et al., 1994, *Gene Therapy* 1: 292-299; Couture et al., 1994, *Hum. Gene Therapy*. 5:, 667-277; and Eiverhand et al., 1995, *Gene Therapy* 2:336-343. Non-viral vectors which are also suitable include naked DNA (see Example Sections 1, 2, 3, and 5), DNA-lipid complexes, for example liposome-mediated or ligand/ poly-L-Lysine conjugates, such as asialoglyco-protein-mediated delivery systems. See for example: Felgner et al., 1994, *J. Biol. Chem.* 269:2550-2561; Derossi et al., 1995, *Restor. Neurol. Neuros*. 8:7-10; and Abcallah et al., 1995, *Biol. Cell* 85:1-7. It is preferred that local cells such as adipose tissue cells or smooth muscle cells, as well as tumor cells, be targeted for delivery and concomitant *in vivo* expression of the respective sVEGF-R protein to promote inhibition of tumor angiogenesis.

A recombinant Ad/sVEGF-RI is a preferred virus for targeting cells proximal to a solid tumor.

An especially preferred recombinant Ad/sVEGF-RI virus is AdHCMVsFLT-1.

Another especially preferred recombinant Ad/sVEGF-RI virus is AdHCMVI1sFLT .

Any membrane bound (mVEGF-R) construct or any FLT-1 or KDR construct encoding a protein deficient in kinase activity may be targeted primarily to vascular endothelial cells in the vicinity of tumor growth. DNA delivery vehicles described above may be utilized to target

any such gene transfer construct to vascular endothelial cells of the mammalian host.

As used herein, "VEGF" or "VEFG-A" refers to vascular endothelial growth factor.

As used herein, "homologue of VEGF" refers to homodimers of VEGF-B, VEGF-C, VEGF-D and PlGF and any functional heterodimers formed between VEGF-A, VEGF-B, VEGF-C, VEGF-D and PlGF, including but not limited to a VEGF-A/PlGF heterodimer.

As used herein, "VEGF-B" refers to vascular endothelial growth factor-B.

As used herein, "VEGF-C" refers to vascular endothelial growth factor-C.

As used herein, "VEGF-D" refers to vascular endothelial growth factor-D.

As used herein, "KDR" or "FLK-1" refers to kinase insert domain-containing receptor or fetal liver kinase.

As used herein, "FLT-1" refers to fms-like tyrosine kinase receptor.

As used herein, "Ad" refers to adenovirus.

As used herein, "HUVEC" refers to human umbilical vein endothelial cell(s).

As used herein, the term "mammalian host" refers to any mammal, including a human being.

As used herein, "sVEGF-R" generically refers to a soluble form of a tyrosine kinase receptor which binds to its respective vascular endothelial growth factor such as VEGF, VEGF-B, VEGF-C, VEGF-D and PlGF without stimulating receptor activation, mitogenesis of vascular endothelial cells or angiogenesis.

As used herein, "sVEGF-RI" or "sFLT-1" refers to the native human soluble form of sFLT, disclosed in U.S. Application Serial No.08/232,538 and presented herein in cDNA form (comprising SEQ ID NO:1) and protein form (SEQ ID NO:2).

As used herein, "VEGF-Rs" refers to a human wild-type VEGF/VEGF homologue specific tyrosine kinase receptor such as FLT-1 and KDR.

As used herein, "mVEGF-R" generically refers to a human wild-type VEGF/VEGF homologue specific tyrosine kinase receptor such which is membrane bound , including but not limited to FLT-1, VEGF-RTMI, KDR, and VEGF-RTMII, as shown in Figure 1.

It is an object of the present invention to provide gene therapy methods to inhibit angiogenesis and growth of solid tumors.

It is also an object of the present invention to utilize a gene or gene fragment of sVEGF-R in gene therapy methods to inhibit angiogenesis and growth of solid tumors.

It is also an object of the present invention to utilize sVEGF-RI in gene therapy methods to inhibit angiogenesis and growth of solid tumors.

It is an object of the present invention to disclose animal models for the determination of efficacy of FLT-1-based constructions for cell delivery and *in vivo* expression in the mammalian host.

It is an object of the present invention to provide recombinant DNA vectors containing sVEGF-RI constructs for use in gene therapy to locally inhibit angiogenesis in a mammalian host.

BRIEF DESCRIPTION OF THE FIGURES

Figure 1 shows a schematic diagram of full length VEGF-Rs (FLT-1 and KDR), the soluble VEGF receptors (sVEGF-RI and sVEGF-RII) and the soluble receptors containing the C-terminal transmembrane region (sVEGF-RTMI and sVEGF-RTMII), with the protein domains of each.

Figure 2 shows the nucleotide sequence of which encodes human sFLT-1 [sVEGF-RI] (SEQ ID NO:1).

Figure 3A and Figure 3B show the amino acid sequence of human sFLT-1 [sVEGF-RI] (SEQ ID NO:2).

Figure 4 shows inhibition of tumor nodules grown in nude mice for HT-1080 mouse cells transiently transfected with pcDNA3/s*flt*-1 (○) or pcDNA3 (●). 3 x 10^6 cells were injected at day 0.

Figure 5 shows inhibition of tumor nodules grown in nude mice for HT-1080 mouse cells stably transfected with pcDNA3/s*flt*-1 (●) or pcDNA3 (○).

Figure 6 shows the survival plot of scid mice injected with (a) D-54MG human glioblastoma cells stably transfected with pcDNA 3 (■); (b) D-54MG human glioblastoma cells stably transfected with pcDNA-s*flt*-1 (□); and (c) untransfected D-54MG human glioblastoma cells (▥).

Figure 7 shows additional data points from the experiment detailed in Figure 6, namely that a CB-17 scid-mouse human-glioma model was used to asses the effect of stable sflt-1 expression on tumor growth and survival. (■)D-54MG human glioblastoma cells stably transfected with pcDNA3; (b) D-54MG human glioblastoma cells stably transfected with pcDNA-s*flt*-1 (▲); and (c) untransfected D-54MG human glioblastoma cells (○).

Figure 8 shows that tumor growth in mice as measured by average volume and mass (±SD) was significantly inhibited by expression of the sflt-1 gene (1-tailed Student's t-test, $p < 0.0001$ for comparison of masses) subcloned into pCDNA1A3, resulting in pcDNAIAsFLT-1.

Figure 9 shows that the tumor masses in mice of sFlt-1 expressing cells were significantly smaller than either the adenovirus treated control cells ($p = 0.035$) or the no virus treated control cells ($p = 0.007$) using the appropriate 1-tailed Student's t-test.

DETAILED DESCRIPTION OF THE INVENTION

The present invention relates to methods of gene therapy for inhibiting VEGF-induced angiogenesis associated with diseases and disorders including, but not limited to, solid tumor growth, tumor metastasis, inflammation, psoriasis, rheumatoid arthritis, hemangiomas, angiofibromas, diabetic retinopathy, and macular

degeneration. These disorders are related in that VEGF acts as a vascular endothelial cell mitogen and chemotactic agent to stimulate local angiogenesis which in turns exacerbates the condition.

The present invention relates to gene transfer of a DNA vector and concomitant *in vivo* expression of a soluble form of a VEGF receptor (sVEGF-R) within the mammalian host which binds VEGF or a VEGF homologue in and around the localized site of the disorder. The formation of a sVEGR-R/VEGF complex will inhibit binding of VEGF to the full-length KDR and FLT-1 tyrosine kinase receptors spanning the vascular endothelial cell surface plasma membrane, thus preventing transduction of the mitogenic and other signals stimulating angiogenesis. In addition, expression of sVEGF-R may also impart a therapeutic effect by binding with membrane associated VEGF full-length receptors to form non-functional receptor heterodimers and thereby inhibit the mitogenic activity of VEGF in a dominant negative manner.

A nucleotide sequence encoding a sVEGF-R for inclusion in a gene therapy vector of the present invention may be chosen from a group of genes encoding tyrosine kinase receptors, namely from the group consisting of *sflt*-1, *flt*-1, KDR (also denoted flk-1), and *flt*-4. The resulting DNA fragment encodes a protein or protein fragment which binds VEGF and inhibits formation of a wild-type, functional VEGF-R/VEGF complex.

A preferred application of the present invention relates to methods inhibiting solid tumor angiogenesis, tumor growth and metastasis by utilizing the disclosed gene therapy methodology. In particular, methods are disclosed for inhibition of primary tumor growth and metastasis by gene transfer of a nucleotide sequence encoding sVEGF-R to a mammalian host. The transferred nucleotide sequence transcribes mRNA and expresses sVEGF-R such that sVEGF-R binds to VEGF in extracellular regions adjacent to the primary tumor and vascular endothelial cells and/or heterodimerizes with full-length VEGF-Rs inhibiting their function. Formation of a sVEGF-R/VEGF-R heterodimeric complexes will prevent VEGF-induced dimerization of

functional full-length VEGF-Rs, antagonizing receptor transphosphorylation-dependent signal transduction associated with vascular endothelial cell-activation and tumor angiogenesis. In addition, expression of sVEGF-R may also impart a therapeutic effect by binding either with or without VEGFs to form non-functional heterodimers with full-length VEGF-Rs and thereby inhibiting the mitogenic and angiogenic activities of VEGFs in constructing the necessary DNA vector. Restriction endonuclease cleavage sites are identified within the receptor DNA and can be utilized directly to excise the extracellular-encoding portion. In addition, PCR techniques as described above may be utilized to produce the desired portion of DNA. It is readily apparent to those skilled in the art that other techniques, which are standard in the art, may be utilized to produce sVEGF-R molecules in a manner analogous to those described above. Such techniques are found, for example, in Maniatis et al. (1982, Molecular Cloning: A Laboratory Manual; Cold Spring Harbor Laboratory, Cold Spring Harbor, New York).

In a particular embodiment of the present invention a DNA fragment encoding a soluble form of the FLT-1 amino acid sequence (see Shibuya, et al., 1990, *Oncogene* 5: 519-524) is utilized in gene therapy protocols. It will be within the purview of the skilled artisan to generate a sVEGF-R construct which binds to VEGF and inhibits forming a complex with wild-type full-length VEGF-R dimers on the cell surface membrane of vascular endothelial cells. Such a construct may be generated by recombinant DNA techniques known in the art using a DNA fragment encoding a partial or complete amino acid sequence of a FLT receptor. Using recombinant DNA techniques, DNA molecules are constructed which encode at least a portion of the VEGF receptor capable of binding VEGF without stimulating mitogenesis or angiogenesis. As described below, *in vivo* delivery of a DNA construct encoding sVEGF-R is targeted to cells and tissue which surround the tumor, including but not limited to vascular endothelial cells, muscle cells, adipose cells, as well as tumor cells and surrounding tissues such as muscle tissue and adipose tissue.

The present invention also relates to therapeutic treatment of the metastatic spread of tumors, the principal cause cancer mortality. Tumor cells can metastasize by entry into the circulatory system, transport to distant sites, implantation back into the surrounding tissue and growth. Inhibition of any step in this process would be expected to inhibit the ultimate establishment and growth of metastatic foci. To this end, an additional aspect of the present invention relates to use of the gene therapy constructs of the present invention, including but not limited to sFlt, to the inhibit the metastatic spread of tumors. The significant inhibition of the establishment of HT1080 metastatic lung foci by sflt expression as shown in Example 2 shows that sflt is effective in inhibiting this process. The sflt-1-transfected HT1080 cell tail vein injection experiment monitors implantation and/or growth of circulating tumor cells, two of the crucial steps in metastatic spread. It is envisioned that sflt may decrease the efficiency of tumor cell extravasation out of blood and into surrounding tissue, possibley by inhibiting VEGF-induced vascular permeability which could facilitate cell migration through vessel walls. Additionally, expression of sFlt is expected to arrest neovascular development within metastatic foci thus diminishing their growth and/or viability.

In another particular embodiment of the present invention a DNA fragment encoding the extracellular ligand binding domain and the transmembrane domain of FLT-1 (see Figure 1) is utilized in gene therapy protocols. Such a DNA construct may be constructed to contain the appropriate wild-type signal sequence such that the proper insertion into the plasma membrane occurs. To this end, it is preferred that viral and non-viral constructs which express VEGF-RTMI (Figure 1) or a biological equivalent thereof, will be targeted substantially to vascular endothelial cells within the region of the tumor.

In another specific embodiment of the present invention, flt-1 is utilized as a template to generate a mutated version of FLT-1 defective in protein kinase activity. A mutant is this class would possess one or more mutations at or around one or more known active sites for protein kinase activity. In other words, the mutant FLT-1 protein will

comprise an extracellular domain, a transmembrane domain, and a mutated intracellular domain. An noted in the previous paragraph regarding delivery of VEGF-RTMI, it is preferred that viral and non-viral constructs which express a mutant FLT-1 be targeted substantially to vascular endothelial cells within the region of the tumor.

An especially preferred template for practicing the present invention is the cDNA encoding a soluble form of FLT-1 (sVEGF-RI), described in Kendall and Thomas (1993, *Proc. Natl. Acad. Sci.* 90: 10705-10709) and U.S. Application Serial No. 08/232,538 which is hereby incorporated by reference. Briefly, a cDNA clone encoding sVEGF-RI was isolated in a two-stage approach employing polymerase chain reaction (PCR) based technology and cDNA library screening. In the first stage, DNA oligonucleotides derived from the extracellular domain sequence information from the known full length FLT, KDR or other VEGF receptor is used to design oligonucleotide primers for the amplification of sVEGF-R-specific DNA fragments. In the second stage, these fragments are cloned to serve as probes for the isolation of complete sVEGF-R cDNA from a commercially available lambda gt10 cDNA library (Clontech) derived from HUVECs (ATCC CRL 1730). This sVEGF-RI cDNA expresses an alternatively spliced form of the FLT-1 precursor mRNA that includes 31 unique amino acid residues at the C-terminal end not found in FLT-1 (see Figure 2 and SEQ ID NO:1 for nucleotide sequence and Figure 3 and SEQ ID NO:2 for amino acid sequence). These 31 unique residues are encoded by a intron that is not removed in this alternatively spliced version. The alternatively spliced mRNA is translated into this intron region until the first stop codon is encountered. This especially preferred template (sflt-1 or sVEGF-RI) for a gene therapy vector will express sVEGF-RI *in vivo* and bind VEGF and/or heterodimerizes with full-length VEGF-Rs (e.g. VEGF-RI/FLT-1 and VEGF-RII/KDR), thus inhibiting tumor angiogenesis.

The cloned sVEGF-RI cDNA obtained through the methods described above may be recombinantly expressed by molecular cloning into an expression vector containing a suitable promoter and oth r appr priate transcription regulatory elements, and transferred into

prokaryotic or eukaryotic host cells to produce recombinant sVEGF-RI. Techniques for such manipulations are fully described in Maniatis, et al.(id.), and are well known in the art.

As noted above, a preferred embodiment of the present invention relates to methods of inhibiting angiogenesis of solid tumors to prevent further tumor growth and eventual metastasis. To this end, any solid tumor or the region surrounding the tumor accessible to gene transfer will be a target for the disclosed therapeutic applications. A sVEGF-R gene or gene fragment, including but not limited to sVEGF-RI and any biologically active truncated version, housed within a recombinant viral- or non-viral-based gene transfer system may be directed to target cells within proximity of the tumor by any number of procedures known in the art, including but not limited to (a) surgical procedures coupled with administration of an effective amount of the DNA to the site in and around the tumor (involving initial removal of a portion or the entire tumor, if possible); (b) injection of the gene transfer vehicle directly into or adjacent to the site of the tumor; and, (c) localized or systemic delivery of the gene transfer vector and/or gene product using techniques known in the art; as listed below.

Therefore, any solid tumor which contains VEGF expressing cells will be a potential target for treatment. Examples, but by no means listed as a limitation, of solid tumors which will be particularly vulnerable to sVEGF-R gene therapy applications are (a) neoplasms of the central nervous system such as, but again not necessarily limited to glioblastomas, astrocytomas, neuroblastomas, meningiomas, ependymomas; (b) cancers of hormone-dependent tissues such as protstate, testicals, uterus, cervix, ovary, mammary carcinomas including but not limited to carcinoma in situ, medullary carcinoma, tubular carcinoma, invasive (infiltrating) carcinomas and mucinous carcinomas; (c) melanomas, including but not limited to cutaneous and ocular melanomas; (d) cancers of the lung which at least include squamous cell carcinoma, spindle carcinoma, small cell carcinoma, adenocarcinoma and large cell carcinoma; and (e) cancers of the gastrointestinal system such as esophageal, stomach, small

intestine, colon, colorectal, rectal and anal region which at least include adenocarcinomas of the large bowel.

Expression vectors are defined herein as DNA sequences that are required for the transcription of cloned copies of genes and the translation of their mRNAs in an appropriate host. Such vectors can be used to express eukaryotic genes in a variety of hosts such as bacteria, bluegreen algae, fungal cells, yeast cells, plant cells, insect cells and animal cells.

Specifically designed vectors allow the shuttling of DNA between hosts such as bacteria-yeast or bacteria-animal or bacteria-insect cells. An appropriately constructed expression vector should contain: an origin of replication for autonomous replication in host cells, selectable markers, a limited number of useful restriction enzyme sites, a potential for high copy number, and active promoters. A promoter is defined as a DNA sequence that directs RNA polymerase to bind to DNA and initiate RNA synthesis. A strong promoter is one which causes mRNAs to be initiated at high frequency. Expression vectors may include, but are not limited to, cloning vectors, modified cloning vectors, specifically designed plasmids or viruses.

A variety of mammalian expression vectors may be used to express recombinant sVEGF-R in mammalian cells. Commercially available mammalian expression vectors which may be suitable for recombinant sVEGF-R expression, include but are not limited to,

Commercially available mammalian expression vectors which may be suitable for recombinant sVEGF-R expression, include but are not limited to, pcDNA3.1 (Invitrogen), pBlueBacHis2 (Invitrogen), pLITMUS28, pLITMUS29, pLITMUS38 and pLITMUS39 (New England Bioloabs), pcDNAI, pcDNAIamp (Invitrogen), pcDNA3 (Invitrogen), pMC1neo (Stratagene), pXT1 (Stratagene), pSG5 (Stratagene), EBO-pSV2-neo (ATCC 37593) pBPV-1(8-2) (ATCC 37110), pdBPV-MMTneo(342-12) (ATCC 37224), pRSVgpt (ATCC 37199), pRSVneo (ATCC 37198), pSV2-dhfr (ATCC 37146), pUCTag (ATCC 37460), and λZD35 (ATCC 37565).

DNA encoding a sVEGF-R, sVEGF-RI or truncated version thereof may also be cloned into an expression vector for expression in a recombinant host cell. Recombinant host cells may be prokaryotic or eukaryotic, including but not limited to bacteria, yeast, mammalian cells including but not limited to cell lines of human, bovine, porcine, monkey and rodent origin, and insect cells including but not limited to drosophila, moth, mosquito and armyworm derived cell lines. The expression vector may be introduced into host cells via any one of a number of techniques including but not limited to transformation, transfection, Ad/polylysine DNA complexes, protoplast fusion, and electroporation. Cell lines derived from mammalian species which may be suitable and which are commercially available, include but are not limited to, CV-1 (ATCC CCL 70), COS-1 (ATCC CRL 1650), COS-7 (ATCC CRL 1651), CHO-K1 (ATCC CCL 61), 3T3 (ATCC CCL 92), NIH/3T3 (ATCC CRL 1658), HeLa (ATCC CCL 2), C127I (ATCC CRL 1616), BS-C-1 (ATCC CCL 26) and MRC-5 (ATCC CCL 171) and HEK 293 cells. Insect cell lines which may be suitable and are commercially available include but are not limited to 3M-S (ATCC CRL 8851) moth (ATCC CCL 80) mosquito (ATCC CCL 194 and 195; ATCC CRL 1660 and 1591) and armyworm (Sf9, ATCC CRL 1711).

A DNA fragment encoding a sVEGF-R, sVEGF-RI or mutant versions thereof may be delivered either systemically or to target cells in the proximity of a solid tumor of the mammalian host by viral or non-viral based methods. Viral vector systems which may be utilized in the present invention include, but are not limited to, (a) adenovirus vectors; (b) retrovirus vectors; (c) adeno-associated virus vectors; (d) herpes simplex virus vectors; (e) SV 40 vectors; (f) polyoma virus vectors; (g) papilloma virus vectors; (h) picarnovirus vectors; and (i) vaccinia virus vectors. Non-viral methods of delivery include but are not necessarily limited to direct injection of naked DNA, such as a recombinant DNA plasmid expression vector described herein comprising a DNA fragment encoding sVEGF-R, VEGF-RTM, or mutated forms of FLT-1 or KDR.

The present invention therefore relates to non-viral recombinant vectors for delivery to the target hosts. To this end, a preferred recombinant plasmid described herein is pcDNA3/sflt-1. An especially preferred recombinant plasmid of the present invention is pcDNAIAsFLT-1, as decribed in Example Section 5.

A recombinant adenovirus (Ad) system is preferred for delivery and prolonged expression within target cells proximal to a solid tumor. A particularly useful adenovirus system used in the present invention is described in Example 4.

A recombinant Ad/sVEGF-RI is a preferred virus for targeting cells proximal to a solid tumor.

An especially preferred recombinant Ad/sVEGF-RI virus is AdHCMVsFLT-1.

Another especially preferred recombinant Ad/sVEGF-RI virus is AdHCMVI1sFLT .

The recombinant Ad/sVEGF-RI viruses of the present invention, including AdHCMVsFLT-1 and AdHCMVI1sFLT, are preferably administered to the host by direct injection into a solid tumor and/or quiescent tissue proximal to the solid tumor, such as adipose or muscle tissue. It will of course be useful to transfect tumor cells in the region of targeted adipose and muscle tissue. Transient expression of a sVEGF-R or VEGF-RTM in these surrounding cells will result in a local extracellular increase in these proteins and will promote binding with VEGF and full-length VEGF-Rs, thus inhibiting formation of activated full-length VEGF-R dimers.

The recombinant Ad/VEGF-RI viruses of the present invention, including AdHCMVsFLT-1 and AdHCMVI1sFLT, may also be delivered by i.v. injection. A recombinant adenovirus delivered by i.v. injection will preferentially infect hepatocytes when administered intravenously, where expression persists for approximately 3-4 weeks subsequent to the initial infection. Suitable titers will depend on a number of factors, such as the particular vector chosen, the host, strength of promoter used and the severity of the disease being treated.

The skilled artisan may alter the titer of virus administered to the patient, depending upon the method of delivery, size of the tumor and efficiency of expression from the recombinant virus. A dose in the range of 10^9 -10^{11} pfu adenovirus is preferred to treat most primary tumors. The skilled artisan will also realize that the number of viral particles encoding the transgene, whether or not replication competent in a complementing host cell, are a relevant dosing unit. In most Adenovirus constructs, there are 50 to 100-fold more DNA containing particles than pfus.

Non-viral vectors which are also suitable include DNA-lipid complexes, for example liposome-mediated or ligand/ poly-L-Lysine conjugates, such as asialoglyco-protein-mediated delivery systems (see, e.g., Felgner et al., 1994, *J. Biol. Chem.* 269: 2550-2561; Derossi et al., 1995, *Restor. Neurol. Neuros.* 8: 7-10; and Abcallah et al., 1995, *Biol. Cell* 85: 1-7).

There are many embodiments of the instant invention which those skilled in the art can appreciate from the specification. To this end, different transcriptional promoters, terminators, carrier vectors or specific gene sequences may be used successfully.

The present invention provides methods of gene therapy which inhibit tumor angiogenesis in a mammalian host. It will be readily apparent to the skilled artisan that various forms of the nucleotide sequence(s) encoding FLT-1, sVEGF-RTMI, sVEGF-R, sVEGF-RI or any mutated version thereof may be utilized to alter the amino acid sequence of the expressed protein. The altered expressed protein may have an altered amino acid sequence, yet still bind to VEGF and in turn inhibit the molecular cascade required to stimulate tumor angiogenesis. For example, various COOH terminal truncated forms of sVEGF-RI are envisioned in the present invention. It will be of ease for the skilled artisan to generate such altered forms upon review of this specification. Any such truncated version of FLT which is soluble and which binds VEGF, a VEGF homologue and/or FLT-1 or KDR is considered a functional equivalent in light of the teachings of this specification. It is also envisioned, as described in the specification, that

membrane bound mutant forms, such as COOH-terminal deletion mutants of FLT-1 and point mutations in the intracellular kinase domain, resulting in a mutant protein substantially defective in protein kinase activity, may be useful as a gene therapy construct for patient delivery and in vivo expression so as to inhibit tumor angiogenesis.

The following examples are provided to illustrate the present invention without, however, limiting the same hereto.

EXAMPLE 1

Isolation of a cDNA Encoding Human sFLT-1

PCR derived products were used as hybridization probes for screening a lambda gt10 cDNA library derived from HUVECs (Clontech). Plating and plaque lifts of the library were performed by standard methods (Maniatis, et al., 1982, Molecular Cloning: A Laboratory Manual; Cold Spring Harbor Laboratory, Cold Spring Harbor, New York). The probes were random-primed labeled with ^{32}P-dCTP to high specific activity and a separate screening of the library (1 x 10^6 plaques per screen) was conducted with each probe. The probes were added to hybridization buffer (50% formamide, 5 x Denhardts, 6 x SSC (1 x SSC = 0.15 M NaCl, 0.015 M Na$_3$citrate·2H$_2$O, pH 7.0), 0.1% SDS, 100 mg/ml salmon sperm DNA) at 1 x 10^6 cpm/ml.

Four positively hybridizing phage were detected using the *flt*-1-specific probe. These positively hybridizing phage were observed to be less than full length *flt*-1.

Two *flt*-1 cDNA clones of about 2.0 kb and 2.7 kb in length were subcloned into pGEM vectors (Promega) and bi-directionally sequenced in their entirety by the chain termination method (Sanger et al., 1977, *Proc. Natl. Acad. Sci* 74: 5463-5467) and shown to contain a single open reading frame of about 569 amino acids. Sequence analysis demonstrated that a portion of the 5' flt-1 coding region was missing from these clones. The remainder of the 5' end was cloned using PCR and combined with the DNA of the clones lacking the 5' end to yield a single op n reading frame encoding about 687 amino acids.

The *flt*-1-derived sVEGF-RI (sflt-1) cDNA nucleotide sequence and deduced amino acid sequence is shown in Figure 2 (nucleotide sequence: SEQ ID NO: 1) and Figure 3 (amino acid sequence: SEQ ID NO: 2). Inspection of the deduced amino acid sequence reveals the presence of a single, large open reading frame of 687 amino acids. By comparison with amino acid sequence of the full length FLT-1 VEGF receptor, 31 amino acids are encoded at the C-terminal end of the sVEGF-RI cDNA which are different from those of FLT-1.

EXAMPLE 2

Inhibition of Tumor Angiogenesis in Mice by Administration of Cells Which Transiently Express sVEGF-RI

The sVEGF-RI cDNA described in Example 1, cloned in pGEM3z and referred to as psflt-1, was digested with BamHI, purified and ligated into BamHI-digested pcDNA3. The resulting plasmid, pcDNA3/sflt-1 (alternatively referred to as SFLT-1), was verified by restriction mapping as well as DNA sequencing of the 5' and 3' 500bp of the BamHI insert. The plasmid was transformed into Top10F' E. coli and purified using Qiagen mega prep and Qiagen Endotoxin removal kit.

The expression plasmid pcDNA3/sflt-1, was mixed with adenovirus-polylysine (AdPl) and transfected into mouse HT-1080 cells (ATCC CRL 1730). Control transfections were performed in identical fashion using unmodified pcDNA3. HT-1080 cells were transfected when 80% confluent and harvested 16-24 hours for subsequent study.

Cell counting on triplicate wells was performed for 3 time points within 7 days of transfection using trypan blue exclusion and revealed no difference in the growth curves between the two groups.

The harvested cells were injected either subcutaneously or via tail vein in nude mice and nodules were measured on selected days for the subcutaneous nodules. For the tail vein injections two sets of experiments were performed. In the first experiment, the animals were sacrificed prior to the development of nodules but there were detectable

differences in the weights of the lungs but the weights were not significantly different. In the second experiment there were definitive differences in the number of nodules per sections with the sflt-1 group having fewer nodules/tissue section.

Transient transfection with pcDNA3/sflt-1 (n=7), as compared to a pcDNA control (n=6), resulted in slower growing tumor nodules on all days examined ($p < 0.01$). These cells had identical growth rates in vitro over a period of 96 hours. Average nodule volumes for sflt-1 transfected cells were 50 mm^3, 75 mm^3 and 190 mm^3 on days 7, 12, and 17, respectively. In contrast, using control pcDNA3 transfected cells, nodules were 151 mm^3 261 mm^3 and 474 mm^3 on days 7, 12 and 17. Similarly, mean lung weights were less in animals receiving pcDNA3/sflt-1 (171 mg, n=3) transfected cells by tail vein injection compared to pcDNA3 controls (205 mg, n=3). Figure 4 shows a marked decrease in tumor volume in nude mice injected with HT-1080 cells which transiently express sVEGF-RI in the form of pcDNA3/sflt-1.

A second study designed to investigate the ability of sFLT-based gene therapy to be applied to treatment of tumor metastasis yielded similar results. HT-1080 cells were transiently transfected with pcDNA3 or pcDNA3/sflt-1. 4×10^6 cells were injected at day 0 via the tail vein of each mouse. The animals were sacrificed after one month and the lungs were extracted, weighed, and examined histologically for tumor burden. Lung histology performed on animals receiving intravenously injected tumors revealed a striking difference between the two groups. pcDNA3 transfected cells were associated with pulmonary intramural tumor spread, massive parenchymal edema and mononuclear infiltrate 20 days after intravenous injection of tumor cells. In contrast, pcDNA3/sflt-1 transfected cells were associated with rare tumor foci, the absence of edema and almost normal lung parenchyma histology. Eight of 9 animals injected with HT-1080 cells transiently expressing sVEGF-RI were clear of tumor growth. Conversely, HT-1080 tumor cells transfected with the pcDNA3 control plasmid showed 2 of 9 without tumor growth while 7 of 9 formed lung

nodules. This data shows that sFLT-based gene therapy applications may be utilized to treat tumor metastasis.

Third, a syngeneic model was examined. Pooled clones were generated for either pcDNA3 or pcDNA3/sflt-1 in GL261 mouse glioma cells. Cell counting of the cells grown in culture revealed no differences between the groups. All 3 pcDNA3 animals grew large tumors after approximately one month. Two of 3 in the sFLT-1 group were tumor free. The third had formed a very small tumor. The histopathology differed but all tumors had a clear malignant appearance.

EXAMPLE 3

Inhibiton of Tumor Angiogenesis in Mice by Administration of Cells Stably Transfected with a cDNA Fragment Encoding sVEGF-RI

The study described in Example 2 was repeated with HT-1080 cells stably transfected with either the pcDNA3 control of pcDNA3/sflt-1. Figure 5 shows a virtually complete inhibition of tumor growth compared to the additional data generated with transiently transfected tumor cells.

To determine the effects of sVEGF-RI on animals survival, the human glioma cell line D-54MG was stably transfected with pcDNA3/psflt-1 or a pcDNA3 control. Clones were pooled and the same number of cells were injected intracranially using a mouse stereotactic device with skull sutures as landmarks. The model has previously been determined to have reliable survival characteristics. Animals were treated identically post-operatively. Figure 6 shows that mice injected with an untransfected control died by day 26, mice injected with pcDNA3 transfected control cells died by day 25, wherein all mice which received pcDNA3/sVEGF-R transfected cells were alive at day 41. Figure 7 shows extended data points from this experiment, showing that the mean survival for D-54MG cells transfected with pcDNA3/sflt-1 was 46.5 days. As noted earlier in this paragraph, D54-MG human glioma cells were transfected with pcDNA3/sflt-1 or pcDNA3 using AdpL transfection.

The cells were subsequently propagated in complete medium containing 400 μg/mL of G418 antibiotic (Gibco BRL, Grand Island, NY) for one month to select for a population of clones that contained the pcDNA3/sflt-1 or pcDNA3 plasmid. The selected cells, representing a population of pooled clones were then harvested using trypsin/EDTA solution (Gibco) and counted using a hemacytometer with trypan blue exclusion. The cells were resuspended to a final concentration of 10^7 cells/100 μL in serum-free DMEM/F12 containing 5% methylcellulose as a vehicle to enhance cell viability. A midline scalp incision was made, followed by a 0.5 mm burr hole 1.5-2.0 mm to the right of the midline and 0.5-1.0 mm posterior to the coronal suture. The cells were loaded into a 100 μL microsyringe and 5 μL was injected sterotactically. A 30 gauge needle mounted on the microsyringe was inserted vertically through the burr hole to a depth of 2.5 mm. Forty-five to sixty seconds after injection, the needle was slowly withdrawn and the incision closed with 9 mm Michel wound clips. Mice were returned to sterile microisolator polycarbonate cages, placed over a heating pad until recovery, and provided autoclaved lab chow and sterile water ad libitum. Animals were assessed twice daily for survival. These results demonstrate that sFLT-1 animals survived longer than historical controls and subsequent controls.

EXAMPLE 4
Construction of AdHCMVsFLT-1

Several systems have been developed for the construction of helper-independent adenovirus (Ad) vectors and have been recently been reviewed by Graham and Prevec (1995, *Mol. Biotech.* 3: 207-220) and Hitt et al. (1995, Techniques for human adenovirus vector construction and characterization, In *Methods in Molecular Genetics, Volume 7. Molecular Virology Techniques Part B*, ed. Kenneth W. Adolph, Academic Press, Inc. Orlando, Florida). All of these systems involve cloning the transgene of interest (coding region flanked by appropriate regulatory sequences) into a shuttle plasmid in which it is flanked by Ad sequences homologous to the region of the viral genome into which the

transgene will be introduced. The DNA from the shuttle plasmid is then rescued into virus by either direct ligation *in vitro* followed by transfection or by *in vivo* homologous recombination following transfection into 293 cells.

E1 shuttle plasmids have been developed for the rescue of inserts into the E1 region. These plasmids contain the left 16% of the Ad genome with a deletion of E1 sequences and cloning sites into which the transgene is introduced. If convenient restriction sites are available in the vector backbone, direct ligation of the shuttle plasmid to purified viral DNA can be performed *in vitro* followed by transfection into 293 cells to generate infectious virus. This method although efficient can require extensive screening if the viral DNA is not completely restricted and in many cases is not practical due to the lack of unique correctly positioned restriction sites. For these reasons many protocols rely on *in vivo* homologous recombination to generate infectious virus. To construct a virus by homologous recombination the shuttle plasmid can be transfected into 293 cells with purified viral DNA that has been restricted in the left end or with viral DNA contained in a second plasmid. As with direct ligation the use of purified viral DNA sometimes requires extensive screening to obtain the desired vector because of the regeneration of parental virus and for this reason plasmid systems are more desirable. A number of plasmid systems have been developed for rescuing inserts into E1 (McGrory et al., 1988, *Virology* 163: 614-6170) or E3 (Ghosh-Choudhury, et al., 1986, *Gene* 50: 161-171; Mittal, et al., 1993,*Virus Res.* 28: 67-90) or both (Bett et al., 1994, *Proc. Natl. Acad. Sci. USA* 91: 8802-8806) regions.

The steps involved in the construction of the helper independent Ad vectors expressing sFLT-1 are outlined below. All steps involve the use of standard protocols for generating adenovirus vectors (Hitt, et al., 1995, In *Methods in Molecular Genetics, Volume 7. Molecular Virology Techniques Part B*, ed. Kenneth W. Adolph, Academic Press, Inc. Orlando, FL.). The coding sequences for sFLT were obtained from plasmid psflt-1 by BamHI digestion and inserted into the BamHI site in the polycloning region of E1 shuttle plasmid

pΔE1sp1HCMV-BGHpA, generating pHCMVsFLT-1. pΔE1sp1HCMV-BGHpA contains Ad5 sequences from bp 1 to 341 and bp 3524 to 5790 with a promoter cassette consisting of the HCMV promoter, a polycloning region and the Bovine growth hormone polyadenylation signal inserted in the E1 anti parallel orientation between Ad5 bp 341 and bp 3524. pHCMVsFLT-1 was then cotransfected into 293 cells with Ad genome plasmid pJM17 (McGrory, et al., 1988, *Virology* 163:614-617) and virus AdHCMVsFLT-1 was generated by *in vivo* recombination between the plasmids. pJM17 contains essentially the entire Ad genome but is non infectious in single transfections of 293 cells since it contains an insertion of a pBR322 derivative at bp 1339 in Ad5 sequences which makes the resulting viral genome too large to package. *In vivo* recombination between pJM17 and pHCMVsFLT-1 generates a vector of a packagable size containing the sFLT-1 expression cassette in the E1 region.

An additional recombinant adenoviral virus is also disclosed. It is essentially the same as the vector described above but utilizes a slightly different HCMV promoter segment consisting of the HCMV promoter and first intron (Intron A). This construct increases expression levels within the mammalian host. To construct this vector sFLT-1 coding sequences were obtained from plasmid pHCMVsFLT-1 (described above) by digestion with KpnI and EcoRI. The sFLT-1 fragment was then inserted into the KpnI and EcoRI sites in E1 shuttle plasmid pHCMVI1-BGHpA, generating pHCMVI1sFLT-1. pHCMVI1sFLT-1 has been cotransfected into 293 cells with Ad genome plasmid pJM17. Alternatively, pHCMVI1sFLT-1 was digested with PacI and ligated with purified viral DNA from the virus AdDE1PacIE3 also digested with PacI. Following the transfection of the ligation products into 293 cells viral plaques were screened to obtain the vector AdHCMVI1sFLT-1.

EXAMPLE 5

Stable Transfection of Human HT1080 Fibrosarcoma Cells with sFlt-1 Inhibits Solid Tumor Growth

Generation of sFlt-1 plasmid- An additional plasmid (pcDNAIAsFLT-1) was constructed that contained the HCMV Intron-A upstream to the sflt-1 cDNA in order to generate HT-1080 clones that secrete increased amounts of sflt-1. This intron has been demonstrated in previous studies to enhance gene expression by 10-100 fold above plasmids containing the HCMV early promoter alone. For the construction of pcDNAIAsFLT-1, pcDNA3 was digested with NruI and KpnI (to remove the HCMV promoter) and ligated with the MscI/KpnI fragment from plasmid pVIJNS-MCS (containing the HCMV promoter and Intron A), generating pcDNAINTA. pcDNAINTA was then digested with KpnI and EcoRI and ligated to a KpnI/EcoRI fragment containing the sFLT-1 coding sequences, generating pcDNAIAsFLT-1.

Selection of HT1080 clones stably transfected with pcDNAIAsFLT-1 and expressing sFlt-1 - Human fibrosarcoma HT1080 tumor cells (Rasheed *et al.*, 1974, *Cancer* 33:1027-1033) were transfected with the plasmid (pcDNAIAsFLT-1) containing the human sFlt-1 gene under the control of the HCMV promoter containing the first HCMV intron and the selectable G418 drug resistance gene. Pooled stably transfected HT1080 cells were plated in 100 cm dishes at a density of 10 and 100 cells/plate. The cells were grown in DMEM supplemented medium [Dulbecco's Modified Eagle Medium/F-12 (DMEM), GIBCOBRL (Cat# 11331-030), 10% fetal bovine serum, (GIBCOBRL Cat# 16000-028) and 1 X penicillin-streptomycin, (GIBCOBRL Cat# 15070-063)] with 500 µg/ml of G418 (GIBCOBRL Cat# 10131-035). The medium was replaced every other day until individual colonies grew to diameters of approximately 2.5 mm. Isolated colonies were treated with trypsin (GIBCOBRL Cat# 25200-056), transferred to 24 well plates and grown to confluence. One ml of medium was removed and tested for VEGF binding activity. The stable clone chosen for further studies had similar growth rates *in vitro* compared to both untransfected cells and cells

transfected with pCDNA3, with cell division occuring approximately every 48 hours.

VEGF Binding Protocol - Heparin-Sepharose CL-6B (Pharmacia Cat# 17-0467-01) was washed 3 times with phosphate buffered saline [PBS] (GIBCOBRL Cat# 20012-027), and resuspended in an equal volume of PBS. One ml of conditioned medium was removed from each well, mixed with 50 µl of the heparin-Sepharose CL-6B slurry and incubated overnight at 4 °C with constant mixing. The heparin-Sepharose beads were pelleted by centrifugation (10,000 x g for 2 min) and washed 3 times with PBS. Bound protein was eluted with 40 µl of PBS containing 1.2 M NaCl. A 10 µl aliquot was removed and added to 10 µl of DMEM/0.2% gelatin, 1 µl of ^{125}I-VEGF (Amersham Cat# IM 274; 100,000 cpm/µl) was added and incubated for 20 min at room temperature. Two µl of 10 mM BS^3 bis(sulfosuccinimidyl) suberate [BS^3], (Pierce Cat# 21579 G) was added to the reaction and incubated for an additional 15 min at room temperature. The crosslinking reaction was stopped by the addition of 20 µl of 2X Laemmli sample buffer (BioRad Cat# 161-0737). Crosslinked complexes were separated by SDS/7.5% PAGE and visualized by autoradiography.

Preparation of selected clones for the tumor growth study - Cells were plated in T-75 flasks and grown to confluence in DMEM supplemented medium. Cells were washed with PBS and trypsinized in 2 ml. Trypsinization was stopped by the addition of 8 ml DMEM supplemented medium and the detached cells were removed and counted. The cells were pelleted by centrifugation (1000 rpm in a Sorvall 6000B table top centrifuge) for 5 min and resuspended in PBS with calcium and magnesium at a final concentration of 1.0 x 10^7 cells/ml and 0.5 ml of cells was injected subcutaneously into mice.

Results - HT-1080 cells (0.5 x 10^6 cells/0.5 ml) stably transfected with either control plasmid or plasmid encoding sflt-1 [pcDNAIAsFLT-1] (n = 10/group) that was cloned and selected for high sflt-1 expression were injected subcutaneously into Balb/c nu/nu female mice (Charles River Laboratories). Tumor length and width were

measured as a function of time and used to calculate tumor volume by the equation:

$$\text{Volume} = 4/3 \cdot p \cdot ((\text{length}/2)(\text{width}/2)(\text{length} + \text{width})/4),$$

which estimates the volume of half a prolate ellipsoid assuming that the height is the average of the length and width. On day 12 after implantation tumor ulceration was visible so the tumors were removed and weighed; expression of sFlt-1 caused a 93% reduction in tumor mass. As shown in Figure 8, tumor growth as measured by average volume and mass (±SD) were significantly inhibited by expression of the sflt-1 gene (1-tailed Student's t-test, $p < 0.0001$ for comparison of masses).

EXAMPLE 6

Infection of Human HT1080 Fibrosarcoma Cells with Replication-Defective Adenovirus Expressing Human sFlt-1 Inhibits Tumor Growth

Generation of sFlt-1 adenoviral constructs are as described in Example Section 4.

Adenoviral infection of HT1080 cells in vitro and implantation in vivo - Cells were plated in T-75 flasks and grown to confluence in DMEM supplemented medium. One flask of cells was trypsinized (2 ml), the cells were removed and resuspended in DMEM supplemented medium and counted to determine the number of cells/plate. Growth medium was removed from flasks and the attached cells were washed with PBS containing calcium and magnesium. Either control adenovirus or adenovirus expressing human sFlt-1 under control of HCMV/intron A were added to flasks at an multiplicity of infection of 20 virus pfu (plaque forming units)/cell in 2 ml of PBS with calcium and magnesium and incubated for 1 hr at 37 °C. The virus was removed and the cells were incubated in a humidified incubator wi 5% CO_2 at 37 °C for and additional 24 hr. Cells were washed with PBS and trypsinized with 2 ml of trypsin. Trypsinization was stopped by the addition of DMEM suppl mented medium and the detached cells were

removed and counted. The cells were pelleted by centrifugation (1000 rpm in a Sorvall 6000B table top centrifuge) for 5 min and resuspended in PBS with calcium and magnesium at a final concentration of 1.0×10^7 cells/ml and 0.5 ml of cells was injected subcutaneously into 6-8 week old Balb/c nu/nu female mice.

Results - Tumor cells that were exposed to either no virus, a control adenovirus or adenovirus expressing sFlt-1 under control of the HCMV/intron A promoter [AdHCMVI1sflt-1] (n = 5/group) were allowed to grow subcutaneously in nude mice. After 11 days of *in vivo* growth the skin over the tumor began to ulcerate in control animals so the tumors were removed from all animals and weighed. The mean group tumor masses ± SEMs are shown in Figure 9.

SEQUENCE LISTING

(1) GENERAL INFORMATION

(i) APPLICANT: MERCK & CO., INC.

(ii) TITLE OF THE INVENTION: GENE THERAPY FOR INHIBITION OF ANGIOGENESIS

(iii) NUMBER OF SEQUENCES: 2

(iv) CORRESPONDENCE ADDRESS:
(A) ADDRESSEE: J. Mark Hand - MERCK & CO., INC.
(B) STREET: 126 EAST LINCOLN AVENUE - P.O. BOX 2000
(C) CITY: RAHWAY
(D) STATE: NJ
(E) COUNTRY: US
(F) ZIP: 07065-0900

(v) COMPUTER READABLE FORM:
(A) MEDIUM TYPE: Diskette
(B) COMPUTER: IBM Compatible
(C) OPERATING SYSTEM: DOS
(D) SOFTWARE: FastSEQ Version 1.5

(vi) CURRENT APPLICATION DATA:
(A) APPLICATION NUMBER:
(B) FILING DATE:
(C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:
(A) APPLICATION NUMBER: U.S. 60/026,641
(B) FILING DATE: September 24, 1996

(viii) ATTORNEY/AGENT INFORMATION:
(A) NAME: Hand, J. Mark
(B) REGISTRATION NUMBER: 36,545
(C) REFERENCE/DOCKET NUMBER: 19810Y

(ix) TELECOMMUNICATION INFORMATION:
(A) TELEPHONE: 908-594-3905
(B) TELEFAX: 908-594-4720
(C) TELEX:

(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 2313 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
(ii) MOLECULE TYPE: cDNA
(iii) HYPOTHETICAL: NO
(iv) ANTISENSE: NO
(v) FRAGMENT TYPE:
(vi) ORIGINAL SOURCE:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

GCGGACACTC CTCTCGGCTC CTCCCCGGCA GCGGCGGCGG CTCGGAGCGG GCTCCGGGGC 60
TCGGGTGCAG CGGCCAGCGG GCCTGGCGGC GAGGATTACC CGGGGAAGTG GTTGTCTCCT 120
GGCTGGAGCC GCGAGACGGG CGCTCAGGGC GCGGGGCCGG CGGCGGCGAA CGAGAGGACG 180
GACTCTGGCG GCCGGGTCGT TGGCCGGGGG AGCGCGGGCA CCGGGCGAGC AGGCCGCGTC 240
GCGCTCACCA TGGTCAGCTA CTGGGACACC GGGGTCCTGC TGTGCGCGCT GCTCAGCTGT 300
CTGCTTCTCA CAGGATCTAG TTCAGGTTCA AAATTAAAAG ATCCTGAACT GAGTTTAAAA 360
GGCACCCAGC ACATCATGCA AGCAGGCCAG ACACTGCATC TCCAATGCAG GGGGGAAGCA 420
GCCCATAAAT GGTCTTTGCC TGAAATGGTG AGTAAGGAAA GCGAAAGGCT GAGCATAACT 480
AAATCTGCCT GTGGAAGAAA TGGCAAACAA TTCTGCAGTA CTTTAACCTT GAACACAGCT 540
CAAGCAAACC ACACTGGCTT CTACAGCTGC AAATATCTAG CTGTACCTAC TTCAAAGAAG 600
AAGGAAACAG AATCTGCAAT CTATATATTT ATTAGTGATA CAGGTAGACC TTTCGTAGAG 660
ATGTACAGTG AAATCCCCGA AATTATACAC ATGACTGAAG GAAGGGAGCT CGTCATTCCC 720
TGCCGGGTTA CGTCACCTAA CATCACTGTT ACTTTAAAAA AGTTTCCACT TGACACTTTG 780
ATCCCTGATG GAAAACGCAT AATCTGGGAC AGTAGAAAGG GCTTCATCAT ATCAAATGCA 840
ACGTACAAAG AAATAGGGCT TCTGACCTGT GAAGCAACAG TCAATGGGCA TTTGTATAAG 900
ACAAACTATC TCACACATCG ACAAACCAAT ACAATCATAG ATGTCCAAAT AAGCACACCA 960
CGCCCAGTCA AATTACTTAG AGGCCATACT CTTGTCCTCA ATTGTACTGC TACCACTCCC 1020
TTGAACACGA GAGTTCAAAT GACCTGGAGT TACCCTGATG AAAAAAATAA GAGAGCTTCC 1080
GTAAGGCGAC GAATTGACCA AAGCAATTCC CATGCCAACA TATTCTACAG TGTTCTTACT 1140
ATTGACAAAA TGCAGAACAA AGACAAAGGA CTTTATACTT GTCGTGTAAG GAGTGGACCA 1200
TCATTCAAAT CTGTTAACAC CTCAGTGCAT ATATATGATA AAGCATTCAT CACTGTGAAA 1260
CATCGAAAAC AGCAGGTGCT TGAAACCGTA GCTGGCAAGC GGTCTTACCG GCTCTCTATG 1320
AAAGTGAAGG CATTTCCCTC GCCGGAAGTT GTATGGTTAA AAGATGGGTT ACCTGCGACT 1380
GAGAAATCTG CTCGCTATTT GACTCGTGGC TACTCGTTAA TTATCAAGGA CGTAACTGAA 1440
GAGGATGCAG GGAATTATAC AATCTTGCTG AGCATAAAAC AGTCAAATGT GTTTAAAAAC 1500
CTCACTGCCA CTCTAATTGT CAATGTGAAA CCCCAGATTT ACGAAAAGGC CGTGTCATCG 1560

TTTCCAGACC CGGCTCTCTA CCCACTGGGC AGCAGACAAA TCCTGACTTG TACCGCATAT 1620
GGTATCCCTC AACCTACAAT CAAGTGGTTC TGGCACCCCT GTAACCATAA TCATTCCGAA 1680
GCAAGGTGTG ACTTTTGTTC CAATAATGAA GAGTCCTTTA TCCTGGATGC TGACAGCAAC 1740
ATGGGAAACA GAATTGAGAG CATCACTCAG CGCATGGCAA TAATAGAAGG AAAGAATAAG 1800
ATGGCTAGCA CCTTGGTTGT GGCTGACTCT AGAATTTCTG GAATCTACAT TTGCATAGCT 1860
TCCAATAAAG TTGGGACTGT GGGAAGAAAC ATAAGCTTTT ATATCACAGA TGTGCCAAAT 1920
GGGTTTCATG TTAACTTGGA AAAAATGCCG ACGGAAGGAG AGGACCTGAA ACTGTCTTGC 1980
ACAGTTAACA AGTTCTTATA CAGAGACGTT ACTTGGATTT TACTGCGGAC AGTTAATAAC 2040
AGAACAATGC ACTACAGTAT TAGCAAGCAA AAAATGGCCA TCACTAAGGA GCACTCCATC 2100
ACTCTTAATC TTACCATCAT GAATGTTTCC CTGCAAGATT CAGGCACCTA TGCCTGCAGA 2160
GCCAGGAATG TATACACAGG GGAAGAAATC CTCCAGAAGA AAGAAATTAC AATCAGAGGT 2220
GAGCACTGCA ACAAAAAGGC TGTTTTCTCT CGGATCTCCA AATTTAAAAG CACAAGGAAT 2280
GATTGTACCA CACAAAGTAA TGTAAAACAT TAA 2313

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 687 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein
(iii) HYPOTHETICAL: NO
(iv) ANTISENSE: NO
(v) FRAGMENT TYPE: internal
(vi) ORIGINAL SOURCE:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

```
Met Val Ser Tyr Trp Asp Thr Gly Val Leu Leu Cys Ala Leu Leu Ser
 1               5                  10                  15
Cys Leu Leu Leu Thr Gly Ser Ser Ser Gly Ser Lys Leu Lys Asp Pro
            20                  25                  30
Glu Leu Ser Leu Lys Gly Thr Gln His Ile Met Gln Ala Gly Gln Thr
        35                  40                  45
Leu His Leu Gln Cys Arg Gly Glu Ala Ala His Lys Trp Ser Leu Pro
    50                  55                  60
Glu Met Val Ser Lys Glu Ser Glu Arg Leu Ser Ile Thr Lys Ser Ala
65                  70                  75                  80
```

```
Cys Gly Arg Asn Gly Lys Gln Phe Cys Ser Thr Leu Thr Leu Asn Thr
                85                  90                  95
Ala Gln Ala Asn His Thr Gly Phe Tyr Ser Cys Lys Tyr Leu Ala Val
            100                 105                 110
Pro Thr Ser Lys Lys Lys Glu Thr Glu Ser Ala Ile Tyr Ile Phe Ile
        115                 120                 125
Ser Asp Thr Gly Arg Pro Phe Val Glu Met Tyr Ser Glu Ile Pro Glu
    130                 135                 140
Ile Ile His Met Thr Glu Gly Arg Glu Leu Val Ile Pro Cys Arg Val
145                 150                 155                 160
Thr Ser Pro Asn Ile Thr Val Thr Leu Lys Lys Phe Pro Leu Asp Thr
                165                 170                 175
Leu Ile Pro Asp Gly Lys Arg Ile Ile Trp Asp Ser Arg Lys Gly Phe
            180                 185                 190
Ile Ile Ser Asn Ala Thr Tyr Lys Glu Ile Gly Leu Leu Thr Cys Glu
        195                 200                 205
Ala Thr Val Asn Gly His Leu Tyr Lys Thr Asn Tyr Leu Thr His Arg
    210                 215                 220
Gln Thr Asn Thr Ile Ile Asp Val Gln Ile Ser Thr Pro Arg Pro Val
225                 230                 235                 240
Lys Leu Leu Arg Gly His Thr Leu Val Leu Asn Cys Thr Ala Thr Thr
                245                 250                 255
Pro Leu Asn Thr Arg Val Gln Met Thr Trp Ser Tyr Pro Asp Glu Lys
            260                 265                 270
Asn Lys Arg Ala Ser Val Arg Arg Arg Ile Asp Gln Ser Asn Ser His
        275                 280                 285
Ala Asn Ile Phe Tyr Ser Val Leu Thr Ile Asp Lys Met Gln Asn Lys
    290                 295                 300
Asp Lys Gly Leu Tyr Thr Cys Arg Val Arg Ser Gly Pro Ser Phe Lys
305                 310                 315                 320
Ser Val Asn Thr Ser Val His Ile Tyr Asp Lys Ala Phe Ile Thr Val
                325                 330                 335
Lys His Arg Lys Gln Gln Val Leu Glu Thr Val Ala Gly Lys Arg Ser
            340                 345                 350
```

```
Tyr Arg Leu Ser Met Lys Val Lys Ala Phe Pro Ser Pro Glu Val Val
        355                 360                 365
Trp Leu Lys Asp Gly Leu Pro Ala Thr Glu Lys Ser Ala Arg Tyr Leu
    370                 375                 380
Thr Arg Gly Tyr Ser Leu Ile Ile Lys Asp Val Thr Glu Glu Asp Ala
385                 390                 395                 400
Gly Asn Tyr Thr Ile Leu Leu Ser Ile Lys Gln Ser Asn Val Phe Lys
                405                 410                 415
Asn Leu Thr Ala Thr Leu Ile Val Asn Val Lys Pro Gln Ile Tyr Glu
            420                 425                 430
Lys Ala Val Ser Ser Phe Pro Asp Pro Ala Leu Tyr Pro Leu Gly Ser
        435                 440                 445
Arg Gln Ile Leu Thr Cys Thr Ala Tyr Gly Ile Pro Gln Pro Thr Ile
    450                 455                 460
Lys Trp Phe Trp His Pro Cys Asn His Asn His Ser Glu Ala Arg Cys
465                 470                 475                 480
Asp Phe Cys Ser Asn Asn Glu Glu Ser Phe Ile Leu Asp Ala Asp Ser
                485                 490                 495
Asn Met Gly Asn Arg Ile Glu Ser Ile Thr Gln Arg Met Ala Ile Ile
            500                 505                 510
Glu Gly Lys Asn Lys Met Ala Ser Thr Leu Val Val Ala Asp Ser Arg
        515                 520                 525
Ile Ser Gly Ile Tyr Ile Cys Ile Ala Ser Asn Lys Val Gly Thr Val
    530                 535                 540
Gly Arg Asn Ile Ser Phe Tyr Ile Thr Asp Val Pro Asn Gly Phe His
545                 550                 555                 560
Val Asn Leu Glu Lys Met Pro Thr Glu Gly Glu Asp Leu Lys Leu Ser
                565                 570                 575
Cys Thr Val Asn Lys Phe Leu Tyr Arg Asp Val Thr Trp Ile Leu Leu
            580                 585                 590
Arg Thr Val Asn Asn Arg Thr Met His Tyr Ser Ile Ser Lys Gln Lys
        595                 600                 605
Met Ala Ile Thr Lys Glu His Ser Ile Thr Leu Asn Leu Thr Ile Met
    610                 615                 620
```

Asn Val Ser Leu Gln Asp Ser Gly Thr Tyr Ala Cys Arg Ala Arg Asn
625 630 635 640
Val Tyr Thr Gly Glu Glu Ile Leu Gln Lys Lys Glu Ile Thr Ile Arg
645 650 655
Gly Glu His Cys Asn Lys Lys Ala Val Phe Ser Arg Ile Ser Lys Phe
660 665 670
Lys Ser Thr Arg Asn Asp Cys Thr Thr Gln Ser Asn Val Lys His
675 680 685

WHAT IS CLAIMED IS:

1. A method of inhibiting angiogenesis of a solid or metastatic tumor in a mammalian host which comprises delivering a DNA vector to said mammalian host, said DNA vector expressing a soluble form of a tyrosine kinase receptor which forms a dimer with VEGF, a VEGF homologue or a VEGF-specific tyrosine kinase receptor protein.

2. The method of claim 1 wherein said mammalian host is a human.

3. The method of claim 2 wherein said DNA vector is a recombinant adenovirus.

4. The method of claim 2 wherein said DNA vector is a recombinant DNA plasmid vector.

5. The method of claim 3 wherein said recombinant adenovirus is delivered by infection into cells a solid tumor or cells adjacent to said solid tumor.

6. The method of claim 4 wherein said cells are selected from the group consisting of adipose cells, muscle cells and vascular endothelial cells.

7. A method of inhibiting solid tumor angiogenesis or metastatic tumor angiogenesis in a mammalian host which comprises delivering a DNA vector to said mammalian host, said DNA vector expressing a soluble form of FLT-1 which forms a dimer with VEGF, a VEGF homologue or a VEGF-specific tyrosine kinase receptor protein.

8. The method of claim 7 wherein said mammalian host is a human.

9. The method of claim 8 wherein said DNA vector is a recombinant adenovirus.

10. The method of claim 8 wherein said DNA vector is a recombinant DNA plasmid vector.

11. The method of claim 9 wherein said recombinant adenovirus is delivered by infection into cells of a solid tumor or cells adjacent to said solid tumor.

12. The method of claim 11 wherein said adjacent escent cells are selected from the group consisting of adipose cells, muscle cells and vascular endothelial cells.

13. The method of claim 12 wherein said recombinant adenovirus AdHCMVsFLT-1.

14. The method of claim 12 wherein said recombinant adenovirus AdHCMVI1sFLT.

15. The method of claim 10 wherein said recombinant DNA plasmid vector is delivered by injection into cells of a solid tumor or cells adjacent to said solid tumor.

16. The method of claim 15 wherein said quiescent cells are selected from the group consisting of adipose cells, muscle cells and vascular endothelial cells.

17. The method of claim 16 wherein said recombinant DNA plasmid vector is pcDNA3/sflt-1.

18. The method of claim 16 wherein said recombinant DNA plasmid vector is pcDNA3IA/sflt-1.

19. A recombinant virus comprising a DNA fragment encoding a soluble form of a VEGF receptor which forms a dimer with VEGF, a VEGF homologue or a VEGF-specific tyrosine kinase receptor protein said recombinant vector containing at least one regulatory sequence which controls expression of said DNA fragment within a mammalian host.

20. A recombinant virus of claim 19 which is a recombinant adenovirus.

21. A recombinant virus of claim 20 wherein said DNA fragment encodes a soluble VEGF receptor, sFLT-1 .

22. A recombinant virus of claim 21 wherein said DNA fragment encodes a human sFLT-1 VEGF receptor as set forth in SEQ ID NO:2.

23. The recombinant virus of claim 22 which is AdHCMVsFLT-1.

24. The method of claim 22 wherein said recombinant adenovirus AdHCMVI1sFLT.

25. A method of determining efficacy of inhibiting tumor angiogenesis, which comprises:

(a) transfecting cultured tumor cells with a DNA vector expressing sFLT;

(b) injecting said transfected tumor cells into a mouse;

(c) sacrificing said mouse after an interval allowing for tumor growth within said mouse; and,

(d) observing formation of tumor nodules is said mouse as compared to a mouse injected with tumor cells transfected with vector along or untransfected tumor cells.

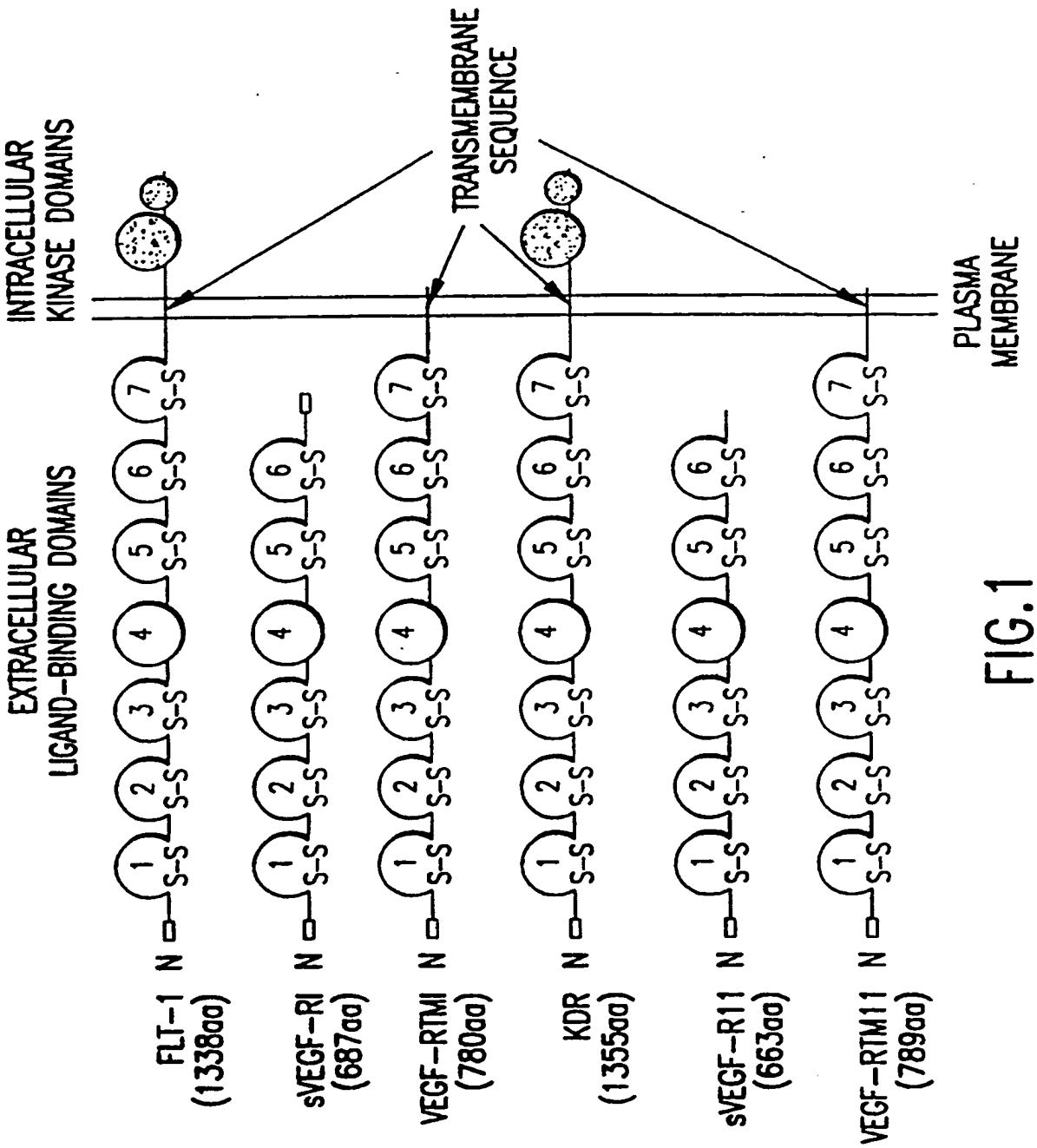
EXTRACELLULAR
LIGAND-BINDING DOMAINS
INTRACELLULAR
KINASE DOMAINS
TRANSMEMBRANE
SEQUENCE
PLASMA
MEMBRANE
FLT-1
(1338aa)
sVEGF-RI
(687aa)
VEGF-RTMI
(780aa)
KDR
(1355aa)
sVEGF-R11
(663aa)
VEGF-RTM11
(789aa)
N
S-S
1
2
3
4
5
6
7

FIG. 1

```
GCGGACACTC CTCTCGGCTC CTCCCCGGCA GCGGCGGCGG CTCGGAGCGG GCTCCGGGGC     60
TCGGGTGCAG CGGCCAGCGG GCCTGGCGGC GAGGATTACC CGGGGAAGTG GTTGTCTCCT    120
GGCTGGAGCC GCGAGACGGG CGCTCAGGGC GCGGGGCCGG CGGCGGCGAA CGAGAGGACG    180
GACTCTGGCG GCCGGGTCGT TGGCCGGGGG AGCGCGGGCA CCGGGCGAGC AGGCCGCGTC    240
GCGCTCACCA TGGTCAGCTA CTGGGACACC GGGGTCCTGC TGTGCGCGCT GCTCAGCTGT    300
CTGCTTCTCA CAGGATCTAG TTCAGGTTCA AAATTAAAAG ATCCTGAACT GAGTTTAAAA    360
GGCACCCAGC ACATCATGCA AGCAGGCCAG ACACTGCATC TCCAATGCAG GGGGGAAGCA    420
GCCCATAAAT GGTCTTTGCC TGAAATGGTG AGTAAGGAAA GCGAAAGGCT GAGCATAACT    480
AAATCTGCCT GTGGAAGAAA TGGCAAACAA TTCTGCAGTA CTTTAACCTT GAACACAGCT    540
CAAGCAAACC ACACTGGCTT CTACAGCTGC AAATATCTAG CTGTACCTAC TTCAAAGAAG    600
AAGGAAACAG AATCTGCAAT CTATATATTT ATTAGTGATA CAGGTAGACC TTTCGTAGAG    660
ATGTACAGTG AAATCCCCGA AATTATACAC ATGACTGAAG GAAGGGAGCT CGTCATTCCC    720
TGCCGGGTTA CGTCACCTAA CATCACTGTT ACTTTAAAAA AGTTTCCACT TGACACTTTG    780
ATCCCTGATG GAAAACGCAT AATCTGGGAC AGTAGAAAGG GCTTCATCAT ATCAAATGCA    840
ACGTACAAAG AAATAGGGCT TCTGACCTGT GAAGCAACAG TCAATGGGCA TTTGTATAAG    900
ACAAACTATC TCACACATCG ACAAACCAAT ACAATCATAG ATGTCCAAAT AAGCACACCA    960
CGCCCAGTCA AATTACTTAG AGGCCATACT CTTGTCCTCA ATTGTACTGC TACCACTCCC   1020
TTGAACACGA GAGTTCAAAT GACCTGGAGT TACCCTGATG AAAAAAATAA GAGAGCTTCC   1080
GTAAGGCGAC GAATTGACCA AAGCAATTCC CATGCCAACA TATTCTACAG TGTTCTTACT   1140
ATTGACAAAA TGCAGAACAA AGACAAAGGA CTTTATACTT GTCGTGTAAG GAGTGGACCA   1200
TCATTCAAAT CTGTTAACAC CTCAGTGCAT ATATATGATA AAGCATTCAT CACTGTGAAA   1260
CATCGAAAAC AGCAGGTGCT TGAAACCGTA GCTGGCAAGC GGTCTTACCG GCTCTCTATG   1320
AAAGTGAAGG CATTTCCCTC GCCGGAAGTT GTATGGTTAA AAGATGGGTT ACCTGCGACT   1380
GAGAAATCTG CTCGCTATTT GACTCGTGGC TACTCGTTAA TTATCAAGGA CGTAACTGAA   1440
GAGGATGCAG GGAATTATAC AATCTTGCTG AGCATAAAAC AGTCAAATGT GTTTAAAAAC   1500
CTCACTGCCA CTCTAATTGT CAATGTGAAA CCCCAGATTT ACGAAAAGGC CGTGTCATCG   1560
TTTCCAGACC CGGCTCTCTA CCCACTGGGC AGCAGACAAA TCCTGACTTG TACCGCATAT   1620
GGTATCCCTC AACCTACAAT CAAGTGGTTC TGGCACCCCT GTAACCATAA TCATTCCGAA   1680
GCAAGGTGTG ACTTTTGTTC CAATAATGAA GAGTCCTTTA TCCTGGATGC TGACAGCAAC   1740
ATGGGAAACA GAATTGAGAG CATCACTCAG CGCATGGCAA TAATAGAAGG AAAGAATAAG   1800
ATGGCTAGCA CCTTGGTTGT GGCTGACTCT AGAATTTCTG GAATCTACAT TTGCATAGCT   1860
TCCAATAAAG TTGGGACTGT GGGAAGAAAC ATAAGCTTTT ATATCACAGA TGTGCCAAAT   1920
GGGTTTCATG TTAACTTGGA AAAAATGCCG ACGGAAGGAG AGGACCTGAA ACTGTCTTGC   1980
ACAGTTAACA AGTTCTTATA CAGAGACGTT ACTTGGATTT TACTGCGGAC AGTTAATAAC   2040
AGAACAATGC ACTACAGTAT TAGCAAGCAA AAAATGGCCA TCACTAAGGA GCACTCCATC   2100
ACTCTTAATC TTACCATCAT GAATGTTTCC CTGCAAGATT CAGGCACCTA TGCCTGCAGA   2160
GCCAGGAATG TATACACAGG GGAAGAAATC CTCCAGAAGA AAGAAATTAC AATCAGAGGT   2220
GAGCACTGCA ACAAAAAGGC TGTTTTCTCT CGGATCTCCA AATTTAAAAG CACAAGGAAT   2280
GATTGTACCA CACAAAGTAA TGTAAAACAT TAA                                2313
```

FIG.2

Met Val Ser Tyr Trp Asp Thr Gly Val Leu Leu Cys Ala Leu Leu Ser
1 5 10 15
Cys Leu Leu Leu Thr Gly Ser Ser Ser Gly Ser Lys Leu Lys Asp Pro
20 25 30
Glu Leu Ser Leu Lys Gly Thr Gln His Ile Met Gln Ala Gly Gln Thr
35 40 45
Leu His Leu Gln Cys Arg Gly Glu Ala Ala His Lys Trp Ser Leu Pro
50 55 60
Glu Met Val Ser Lys Glu Ser Glu Arg Leu Ser Ile Thr Lys Ser Ala
65 70 75 80
Cys Gly Arg Asn Gly Lys Gln Phe Cys Ser Thr Leu Thr Leu Asn Thr
85 90 95
Ala Gln Ala Asn His Thr Gly Phe Tyr Ser Cys Lys Tyr Leu Ala Val
100 105 110
Pro Thr Ser Lys Lys Lys Glu Thr Glu Ser Ala Ile Tyr Ile Phe Ile
115 120 125
Ser Asp Thr Gly Arg Pro Phe Val Glu Met Tyr Ser Glu Ile Pro Glu
130 135 140
Ile Ile His Met Thr Glu Gly Arg Glu Leu Val Ile Pro Cys Arg Val
145 150 155 160
Thr Ser Pro Asn Ile Thr Val Thr Leu Lys Lys Phe Pro Leu Asp Thr
165 170 175
Leu Ile Pro Asp Gly Lys Arg Ile Ile Trp Asp Ser Arg Lys Gly Phe
180 185 190
Ile Ile Ser Asn Ala Thr Tyr Lys Glu Ile Gly Leu Leu Thr Cys Glu
195 200 205
Ala Thr Val Asn Gly His Leu Tyr Lys Thr Asn Tyr Leu Thr His Arg
210 215 220
Gln Thr Asn Thr Ile Ile Asp Val Gln Ile Ser Thr Pro Arg Pro Val
225 230 235 240
Lys Leu Leu Arg Gly His Thr Leu Val Leu Asn Cys Thr Ala Thr Thr
245 250 255
Pro Leu Asn Thr Arg Val Gln Met Thr Trp Ser Tyr Pro Asp Glu Lys
260 265 270
Asn Lys Arg Ala Ser Val Arg Arg Arg Ile Asp Gln Ser Asn Ser His
275 280 285
Ala Asn Ile Phe Tyr Ser Val Leu Thr Ile Asp Lys Met Gln Asn Lys
290 295 300
Asp Lys Gly Leu Tyr Thr Cys Arg Val Arg Ser Gly Pro Ser Phe Lys
305 310 315 320
Ser Val Asn Thr Ser Val His Ile Tyr Asp Lys Ala Phe Ile Thr Val
325 330 335
Lys His Arg Lys Gln Gln Val Leu Glu Thr Val Ala Gly Lys Arg Ser
340 345 350
Tyr Arg Leu Ser Met Lys Val Lys Ala Phe Pro Ser Pro Glu Val Val
355 360 365
Trp Leu Lys Asp Gly Leu Pro Ala Thr Glu Lys Ser Ala Arg Tyr Leu
370 375 380
Thr Arg Gly Tyr Ser Leu Ile Ile Lys Asp Val Thr Glu Glu Asp Ala
385 390 395 400
Gly Asn Tyr Thr Ile Leu Leu Ser Ile Lys Gln Ser Asn Val Phe Lys
405 410 415
Asn Leu Thr Ala Thr Leu Ile Val Asn Val Lys Pro Gln Ile Tyr Glu
420 425 430
Lys Ala Val Ser Ser Phe Pro Asp Pro Ala Leu Tyr Pro Leu Gly Ser
435 440 445
Arg Gln Ile Leu Thr Cys Thr Ala Tyr Gly Ile Pro Gln Pro Thr Ile
450 455 460
Lys Trp Phe Trp His Pro Cys Asn His Asn His Ser Glu Ala Arg Cys
465 470 475 480
Asp Phe Cys Ser Asn Asn Glu Glu Ser Phe Ile Leu Asp Ala Asp Ser
485 490 495

FIG. 3A

Asn Met Gly Asn Arg Ile Glu Ser Ile Thr Gln Arg Met Ala Ile Ile
500 505 510
Glu Gly Lys Asn Lys Met Ala Ser Thr Leu Val Val Ala Asp Ser Arg
515 520 525
Ile Ser Gly Ile Tyr Ile Cys Ile Ala Ser Asn Lys Val Gly Thr Val
530 535 540
Gly Arg Asn Ile Ser Phe Tyr Ile Thr Asp Val Pro Asn Gly Phe His
545 550 555 560
Val Asn Leu Glu Lys Met Pro Thr Glu Gly Glu Asp Leu Lys Leu Ser
565 570 575
Cys Thr Val Asn Lys Phe Leu Tyr Arg Asp Val Thr Trp Ile Leu Leu
580 585 590
Arg Thr Val Asn Asn Arg Thr Met His Tyr Ser Ile Ser Lys Gln Lys
595 600 605
Met Ala Ile Thr Lys Glu His Ser Ile Thr Leu Asn Leu Thr Ile Met
610 615 620
Asn Val Ser Leu Gln Asp Ser Gly Thr Tyr Ala Cys Arg Ala Arg Asn
625 630 635 640
Val Tyr Thr Gly Glu Glu Ile Leu Gln Lys Lys Glu Ile Thr Ile Arg
645 650 655
Gly Glu His Cys Asn Lys Lys Ala Val Phe Ser Arg Ile Ser Lys Phe
660 665 670
Lys Ser Thr Arg Asn Asp Cys Thr Thr Gln Ser Asn Val Lys His
675 680 685

FIG. 3B

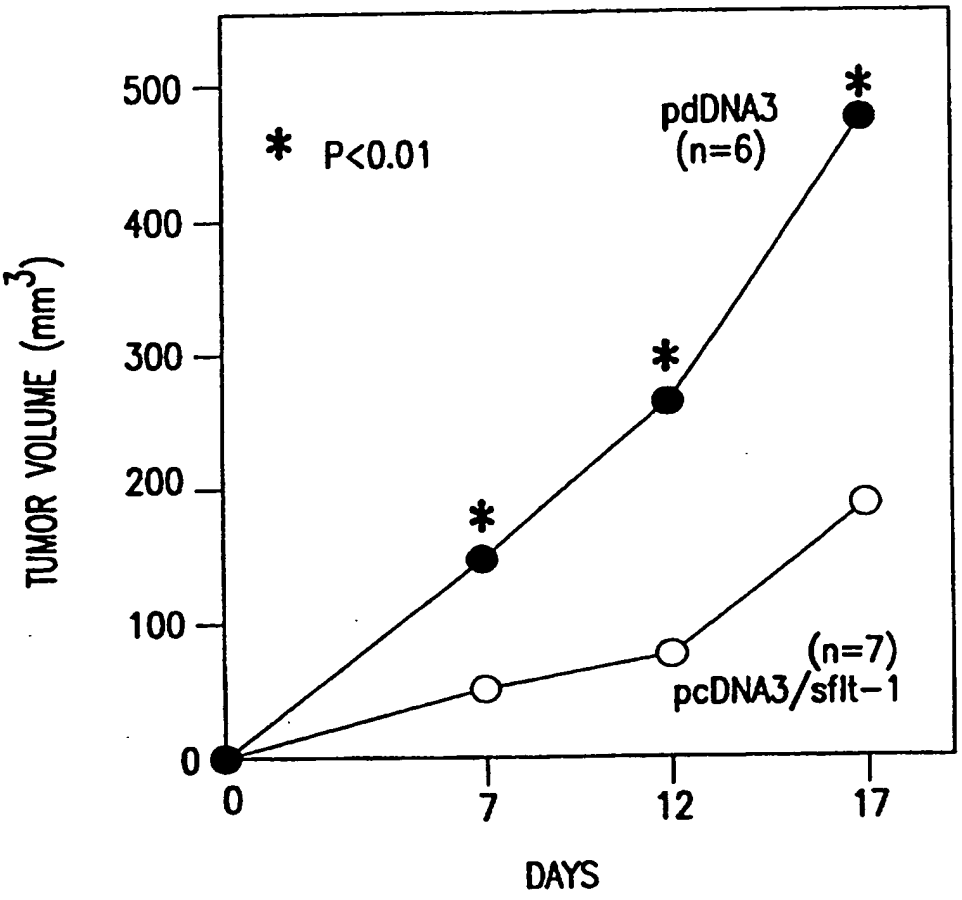
500
400
300
200
100
0
TUMOR VOLUME (mm3)
* P<0.01
pdDNA3
(n=6)
(n=7)
pcDNA3/sflt-1
0
7
12
17
DAYS

FIG.4

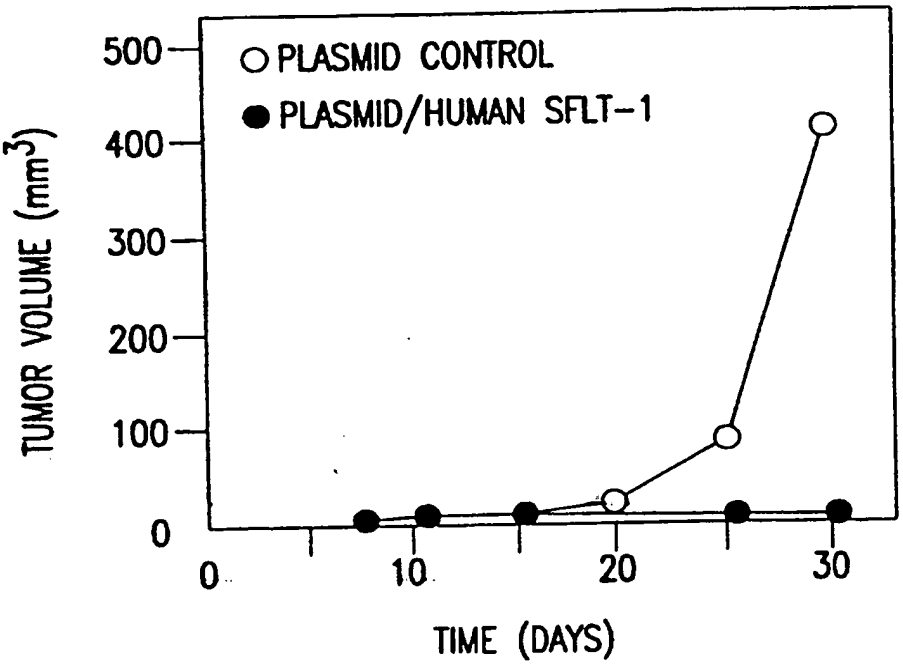
PLASMID CONTROL
PLASMID/HUMAN SFLT-1
TUMOR VOLUME (mm3)
500
400
300
200
100
0
0
10
20
30
TIME (DAYS)

FIG.5

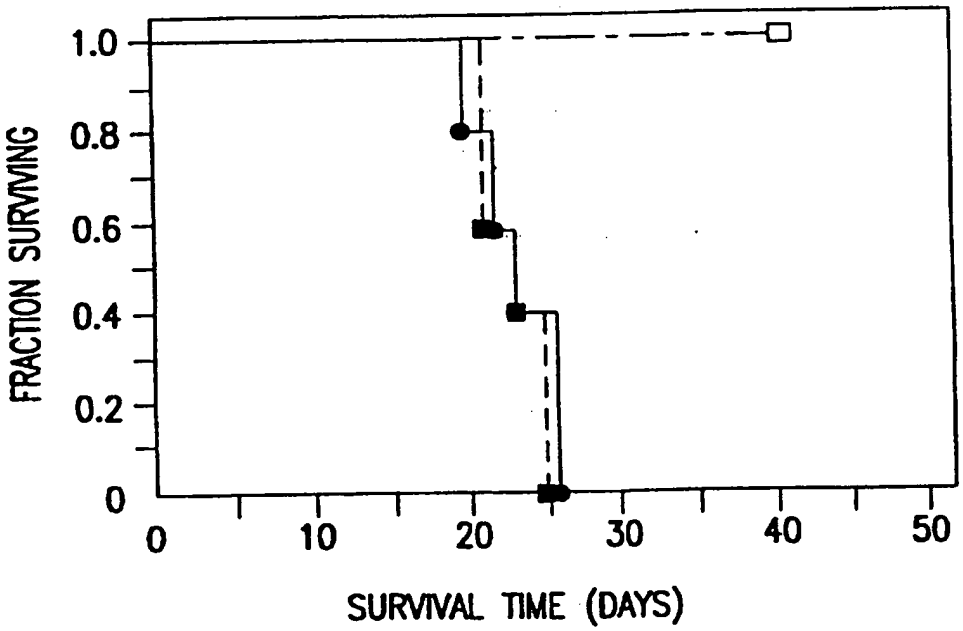
FRACTION SURVIVING
1.0
0.8
0.6
0.4
0.2
0
0
10
20
30
40
50
SURVIVAL TIME (DAYS)

FIG.6

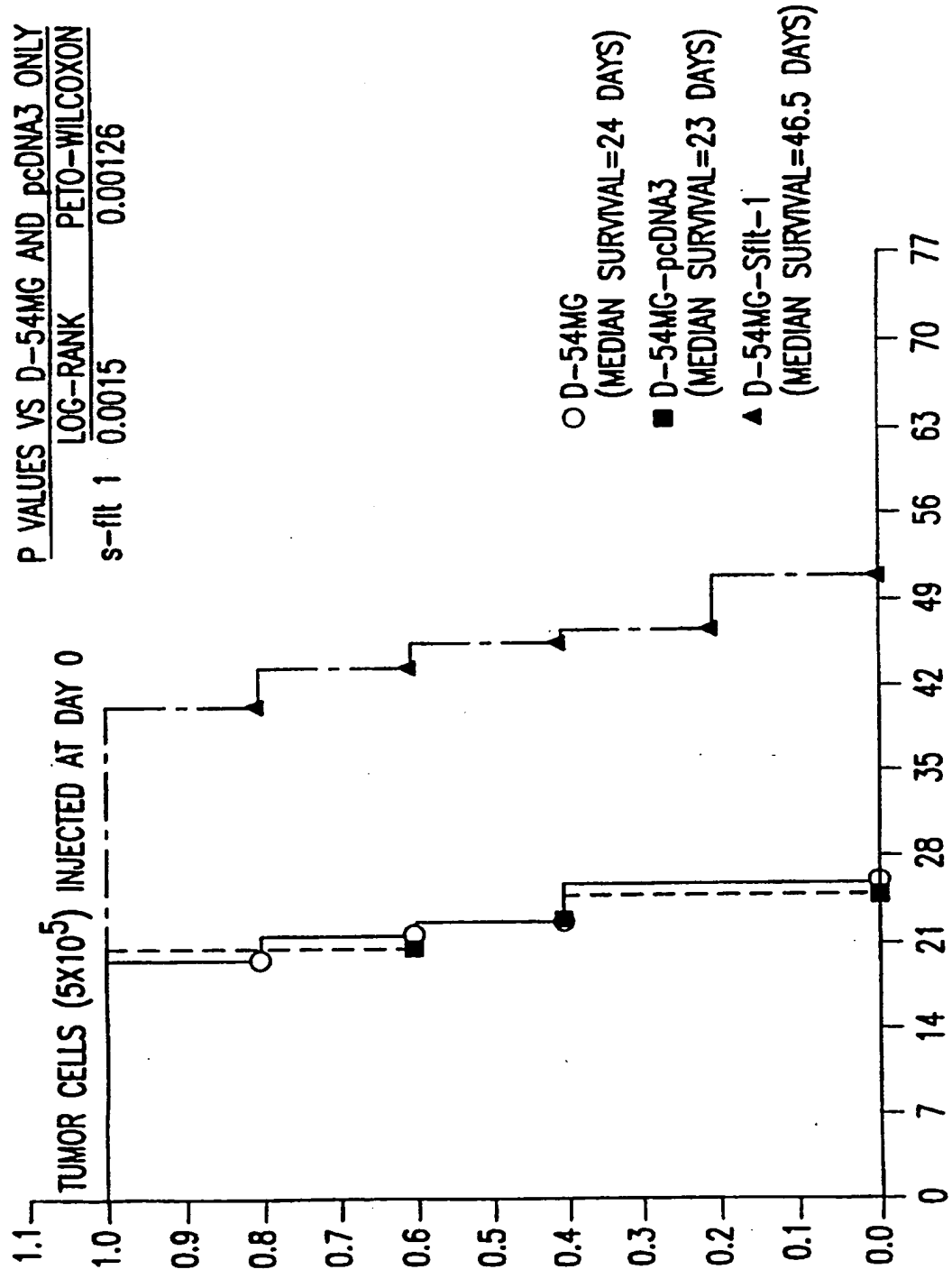
P VALUES VS D-54MG AND pcDNA3 ONLY
LOG-RANK PETO-WILCOXON
s-flt 1 0.0015 0.00126
TUMOR CELLS (5X10^5) INJECTED AT DAY 0
○ D-54MG (MEDIAN SURVIVAL=24 DAYS)
■ D-54MG-pcDNA3 (MEDIAN SURVIVAL=23 DAYS)
▲ D-54MG-Sflt-1 (MEDIAN SURVIVAL=46.5 DAYS)
PROPORTION SURVIVING
SURVIVAL AFTER TUMOR INDUCTION (DAYS)
1.1
1.0
0.9
0.8
0.7
0.6
0.5
0.4
0.3
0.2
0.1
0.0
0
7
14
21
28
35
42
49
56
63
70
77

FIG.7

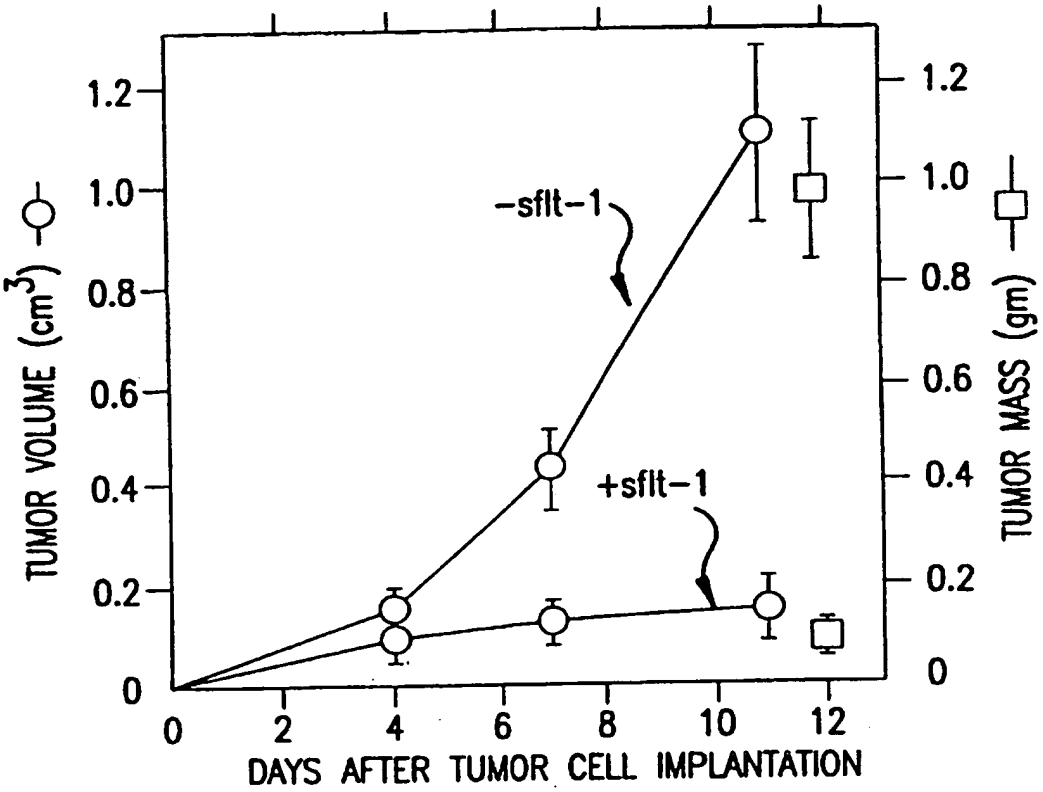
TUMOR VOLUME (cm^3)
TUMOR MASS (gm)
-sflt-1
+sflt-1
0
0.2
0.4
0.6
0.8
1.0
1.2
0
2
4
6
8
10
12
DAYS AFTER TUMOR CELL IMPLANTATION

FIG.8

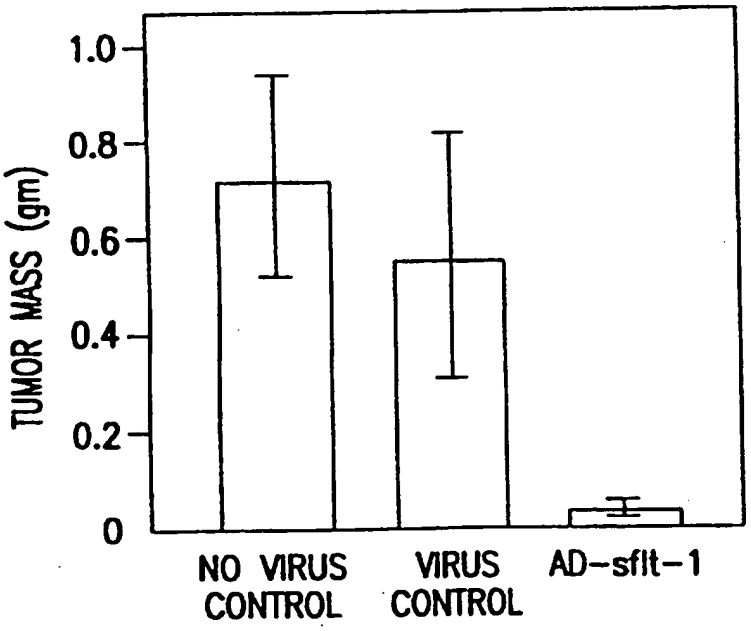
TUMOR MASS (gm)
0
0.2
0.4
0.6
0.8
1.0
NO VIRUS CONTROL
VIRUS CONTROL
AD-sflt-1

FIG.9

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US97/17044

A. CLASSIFICATION OF SUBJECT MATTER

IPC(6) :A61K 48/00, 49/00; C12Q 1/68, 1/70; C12N 15/85, 15/86
US CL :424/93.2, 93.6; 435/5, 6, 320.1
According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 424/93.2, 93.6; 435/5, 6, 320.1

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

APS, Dialog, Biosis, Medline, Biotech
Search terms: gene therapy, adenovirus, angiogenesis, tyrosine kinase receptor, VEGF, FLT-1

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	US 5,194,596 A (TISCHER et al.) 16 March 1993, columns 1-2.	1-6, 19-23
A	MILLAUER et al. Glioblastoma Growth Inhibited in vivo by a Dominant-Negative Flk-1 Mutant. Nature. 10 February 1994, Vol. 367, pages 576-579, especially pages 577-578.	1-25
A	AIELLO et al. Suppression of Retinal Neovascularization in vivo by Inhibition of Vascular Endothelial Growth Factor (VEGF) Using Soluble VEGF-Receptor Chimeric Proteins. Proceedings of the National Academy of Science, USA. November 1995, Vol. 92, pages 10457-10461, especially page 10460.	1-18

[X] Further documents are listed in the continuation of Box C. [ ] See patent family annex.

*	Special categories of cited documents:	"T"	later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
"A"	document defining the general state of the art which is not considered to be of particular relevance	"X"	document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
"E"	earlier document published on or after the international filing date	"Y"	document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
"L"	document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	"&"	document member of the same patent family
"O"	document referring to an oral disclosure, use, exhibition or other means		
"P"	document published prior to the international filing date but later than the priority date claimed		

Date of the actual completion of the international search 22 NOVEMBER 1997	Date of mailing of the international search report 23 DEC 1997
Name and mailing address of the ISA/US Commissioner of Patents and Trademarks Box PCT Washington, D.C. 20231 Facsimile No. (703) 305-3230	Authorized officer DAVID GUZO Telephone No. (703) 308-0196

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INTERNATIONAL SEARCH REPORT

International application No.
PCT/US97/17044

C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	KENDALL et al. Inhibition of Vascular Endothelial Cell Growth Factor Activity by an Endogenously Encoded Soluble Receptor. Proceedings of the National Academy of Science, USA. Vol. 90, pages 10705-10709, especially 10708-10709.	1-25